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# Sleep Quality, Nutrient Intake, and Social Development Index Predict Metabolic Syndrome in the Tlalpan 2020 Cohort: A Machine Learning and Synthetic Data Study

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**Abstract:** This study investigated the relationship between Metabolic Syndrome (MetS), sleep disorders, the consumption of some nutrients, and social development factors, focusing on gender differences in an unbalanced dataset from a Mexico City cohort. We used data balancing techniques like SMOTE and ADASYN after employing machine learning models like random forest and RPART to predict MetS. Random forest excelled, achieving significant, balanced accuracy, indicating its robustness in predicting MetS and achieving a balanced accuracy of approximately 87%. Key predictors for men included body mass index and family history of gout, while waist circumference and glucose levels were most significant for women. In relation to diet, sleep quality, and social development, metabolic syndrome in men was associated with high lactose and carbohydrate intake, educational lag, living with a partner without marrying, and lack of durable goods, whereas in women, best predictors in these dimensions include protein, fructose, and cholesterol intake, copper metabolites, snoring, sobbing, drowsiness, sanitary adequacy, and anxiety. These findings underscore the need for personalized approaches in managing MetS and point to a promising direction for future research into the interplay between social factors, sleep disorders, and metabolic health, which mainly depend on nutrient consumption by region.

**Keywords:** poor quality sleep; social development index; nutrients; machine learning; features selection; balancing methods; Mexico City; Tlalpan 2020 cohort

# 1. Introduction

Metabolic Syndrome (MetS) is a condition that increases the risk of developing or worsening several serious health conditions such as diabetes, heart disease, and stroke, as well as cognitive decline and dementia [1]. Sleep disturbances such as insomnia, apnea, and snoring, linked to MetS, can exacerbate these health risks [2,3]. In 2017, the National Health and Nutrition Survey of Mexico [4] estimated the prevalence of sleep disorders in



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**Copyright:** © 2024 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). Mexicans using a sample of 8649 people older than 18 years old. The results showed a prevalence of snoring while sleeping of 48.5%, difficulty sleeping of 36.9%, and tiredness or fatigue during the day of 32.4%; likewise, insomnia was 18.8% more prevalent in women. Regarding apnea, the results indicated that 23.7% had a higher risk of presenting apnea, especially the populations of those who were overweight and obese, hypertensive, and those over 40 years of age. In another study [5], the prevalence of insomnia was 36.7%, being more common among women (with a prevalence of 41.9%) than men (with a prevalence of 36.7%). Effective treatment for sleep disorders hinges on identifying their specific type and underlying causes, highlighting the ongoing need for improved diagnosis and treatment strategies.

The prevalence data on sleep disorders underscore the importance of understanding their impact on conditions like MetS. This underscores the necessity of employing tools such as the Medical Outcomes Study Sleep Scale (MOS) [6] in research to assess sleep quality and its influence on health. Its widespread use in diverse research studies [7–9] has deepened studies of how sleep disorders affect various health conditions and populations, thanks to its ability to measure multiple sleep-related aspects.

Similarly, nutrition and specific nutrients play crucial roles in developing and managing MetS [10]. MetS is a cluster of conditions that includes abdominal obesity, insulin resistance, dyslipidemia, and hypertension. Poor dietary choices and other lifestyle factors can contribute to developing and exacerbating these risk factors [11,12]. Excessive caloric intake, especially from high-fat and high-sugar diets, contributes to obesity; in consequence, it can contribute to insulin resistance, which is a key feature of metabolic syndrome. Low consumption of dietary fiber, commonly found in fruits, vegetables, and whole grains, is associated with insulin resistance. Diets high in saturated and trans fats can lead to dyslipidemia, which is characterized by elevated levels of triglycerides and low-density lipoprotein cholesterol and decreased high-density lipoprotein cholesterol. This lipid profile is a risk factor for cardiovascular diseases associated with metabolic syndrome. In contrast, omega-3 fatty acids, found in fatty fish, flax seeds, and walnuts, have been associated with favorable lipid profiles and may have a protective effect against metabolic syndrome [13–16]. As expected, nutrition and dietary habits are associated with MetS; various research has found the contributions of nutrients through applying diverse statistical models on the increasing or decreasing risk [17–19].

In the same way, another factor significantly associated with MetS is the social development index (SDI) [20], which is a composite measure of social and economic development. The SDI serves as a metric to evaluate the well-being and social progress in Mexico. Originating in the early 2000s and modeled after the Human Development Index (HDI), the SDI categorizes the level of social development in territorial units. These units correspond, for instance, to the subdivision of municipal geostatistical areas in Mexico City. The SDI employs a methodology established by the National Council for the Evaluation of Social Development Policy (Consejo Nacional de Evaluación de la Política de Desarrollo Social, CONEVAL) for its calculation (refer to Methods for further details on the SDI) [21].

Countries with higher SDI scores tend to have better health outcomes, including lower rates of MetS [22], and an additional study connects the risk of MetS with economic and social vulnerability as well as inappropriate nutrition profiles [23]. The evidence suggests a close association between the SDI and sleep disturbances, which is a relationship influenced by socioeconomic factors such as income level and education. These factors directly affect access to health services and lifestyle habits, such as diet and physical activity, which are essential for maintaining optimal sleep quality. Analyzing how the SDI and sleep disturbances interact with MetS is crucial to unravel the social and economic determinants that shape these complex interconnections. Understanding these dynamics will not only facilitate the identification of the types of sleep disorders that increase the prevalence of MetS but will also contribute to developing more effective strategies for its prevention and treatment, thus improving overall health and well-being. For this reason, developing

automated methods for diagnosing sleep disorders, identifying the determinants of the SDI, and predicting MetS have become fields of significant research interest.

In the case of sleep disruption, machine learning has shown promise in improving the accuracy and efficiency of the diagnosis process. The work of Mencar et al. [24] presents the application of five machine learning models to predict the severity of obstructive sleep apnea syndrome (OSAS) using polysomnography data, where the random forest model obtained the highest accuracy (90.91%) and relevant features such as respiratory rate and oxygen saturation were extracted. Another study [25] applies a machine learning model to predict the presence of OSAS using clinical and demographic data. The random forest model performed best, achieving an accuracy of 87.1%. The most important predictors were body mass index (BMI), age, and gender, as well as additional predictors such as neck circumference and smoking.

In another study by Eyvazlou et al. [26], an ANN model was developed to predict MetS based on sleep quality and work-related risk factors. The results showed that the ANN model could identify individuals at risk of MetS with a sensitivity of 74.1% and a specificity of 76.2%. Moreover, other studies [27,28] have also applied machine learning to understand the social determinants that affect and influence the health of individuals.

However, despite the excellent results described in previous studies, one of the most common challenges in medical diagnoses is the issue of class imbalance. This problem significantly impacts the performance of classifiers, as they tend to exhibit a bias towards the majority class, resulting in skewed outcomes. In this context, authors such as Kim et al. [29] propose a prediction model that utilizes balancing techniques to identify middle-aged Korean individuals at a high risk of MetS. The dataset used in their study comprises age, gender, anthropometric data, sleep quality, and blood indicators of 1991 individuals. The results showed that XGBoost (using Scikit-learn library in Python ver. 3.8.5), employing SMOTE, achieved an AUC of 85.1%.

The present study examines the connection between the SDI, sleep disturbances, types of nutrients consumed, and MetS within a cohort from Mexico City. We aim to identify critical factors that may be key to reducing MetS incidence or severity by applying machine learning algorithms. Additionally, we will use data balancing techniques to improve the predictive performance of our models and enhance feature selection. By incorporating these methods, we aim to uncover valuable insights and contribute to developing more accurate and practical approaches for addressing MetS.

### 2. Materials and Methods

# 2.1. Data

Data for this study were derived from the baseline assessment of a cohort called Tlalpan 2020 from the National Institute of Cardiology Ignacio Chávez in Mexico City [30]. This project was authorized by the Institutional Ethics Committee of the National Institute of Cardiology Ignacio Chavez under code 13-802. The dataset used in this investigation includes data from 3156 volunteers (all of them were informed of the research purposes and signed a letter of informed consent) about their anthropometric measurements, consumption of alcohol and tobacco, level of physical activity, level of economic income, level of education, anxiety, family history health, biomedical evaluation, quality of sleep, and the amount of nutrients consumed.

### 2.1.1. Quality of Sleep

The sleep quality was measured using MOS [6], a self-report for assessing sleep quality and quantity. This questionnaire includes 12 items about sleep disruption, snoring, sleep shortness of breath or headache, sleep adequacy, and sleep somnolence; it additionally measures the number of hours of sleep per day over the previous four weeks. The MOS has been used in several studies, such as discriminating the quality of sleep among a Spanish postmenopausal population [9], diagnosing cases of apnea [7,8], and identifying sleep disturbance in patients with rheumatoid arthritis [31], among others.

### 2.1.2. Clinical and Anthropometric Parameters

Clinical and anthropometric data such as systolic blood pressure (SBP) and diastolic blood pressure (DBP) (measured according to standard procedure [32]) were collected, as well as waist circumference (WC), height and weight (measured according to ISAK [33]) for calculation BMI, and the height–waist index (WHtR). These were calculated from primary measurement data.

# 2.1.3. Biochemical Evaluation

The following laboratory test measurements corresponding to blood samples were included: glucose (GLU), triglycerides (TRIG), HDL cholesterol (HDL), LDL cholesterol (LDL), uric acid (URIC), atherogenic index (IAT), and sodium (NA).

# 2.1.4. Social Development Index

Comprising key dimensions associated with education, health, and housing, the SDI incorporates specific indicators for the evaluation of each dimension. The weight assigned to each indicator varies based on its significance in the overall assessment of social development. The resulting scores are aggregated to yield a score for each dimension. The SDI value facilitates the ordering of territorial units based on their achieved levels of development, classified as Very Low, Low, Medium, and High [34,35].

The SDI indicators (as reported in reference [21]) are briefly described below:

- Quality and available space in the home (QUA\_HOUS): The quality of housing is measured by the type of flooring, and the amount of living space is indicated by the number of people per bedroom, with two being the standard.
- Educational access (EDULAG): This indicator measures the proportion of people aged 18–59 who have completed secondary school or have received 13 years of schooling, which is considered a minimum standard for well-being.
- Access to social security and/or Medical Service (HEALTHAC): This indicator measures the coverage of any of the Mexican health systems.
- Durable goods (DURAB): This indicator measures possession of material goods whose value is equal to or greater than USD 17.81, or possession of at least three items such as a television, gas stove, computer, refrigerator, or washing machine.
- Sanitary adequacy (SANITRY): This indicator measures the availability of a water supply, toilet facilities, and access to a drainage system.
- Electricity access (ENER\_AD): This indicator measures whether or not there is adequate access to electricity.

### 2.1.5. Habits and Factors Associated with Lifestyle

Furthermore, habit data were also collected, such as habitual smoking, alcohol consumption, and physical activity (calculated based on the International Physical Activity Questionnaire, IPAQ, Ref. [36] by metabolic equivalent minutes/week, which are classified in the following categories: low, moderate, and high).

Education level was collected and classified into three categories: primary school, high school, and university studies, as well as postgraduate school. Similarly, we collected the level of economic income, which was classified into three categories based on the Mexican peso income paid monthly: low (MXN 1.00 to MXN 6600.00), medium (MXN 6601.00 to MXN MXN 11,000.00), and high (more than MXN11,000.00).

### 2.1.6. Psychological Stress Level

We used the State-Trait Anxiety Inventory (STAI) to collect data about psychological stress levels, which were categorized into five categories: high (>65), moderate (56–65), medium (46–55), minor (36–45), and low (<35) [37,38].

To gather information about the frequency of food consumption and other dietary products, we utilized a software tool called the "Evaluation of Nutritional Habits and Nutrient Consumption System" from the National Institute of Public Health [39]. This system examines the meals individuals have consumed over a day within the previous year and computes the quantity of nutrients ingested.

All data mentioned in this section are presented in the Table 1.

Table 1. Dataset variables.

AGEageContinuousWEIGHTweightContinuousHEIGHTheightContinuousBMIbody mass indexContinuousWCwaistContinuousSBPsystolic blood pressureContinuousDBPdiastolic blood pressureContinuousLIV_TOGcommon-law marriageDichotomousSINGLEsingleDichotomousDIVORCdivorcedDichotomousVALUEsocial development index by valueContinuousSTRATUMsocioeconomic stratumContinuousQUA_HOUSquality and living spaceContinuousHEALTHACaccess to healthcare and social securityContinuousDILRABdurable goodsContinuous	Name Variable	Description	Туре
WEIGHTweightContinuousHEIGHTheightContinuousBMIbody mass indexContinuousWCwaistContinuousSBPsystolic blood pressureContinuousDBPdiastolic blood pressureContinuousLIV_TOGcommon-law marriageDichotomousMARRIEDmarriedDichotomousSINGLEsingleDichotomousDIVORCdivorcedDichotomousVALUEsocial development index by valueContinuousSTRATUMsocioeconomic stratumContinuousQUA_HOUSquality and living spaceContinuousHEALTHACaccess to healthcare and social securityContinuousDILRABdurable goodsContinuous	AGE	age	Continuous
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WCwaistContinuousSBPsystolic blood pressureContinuousDBPdiastolic blood pressureContinuousLIV_TOGcommon-law marriageDichotomousMARRIEDmarriedDichotomousSINGLEsingleDichotomousDIVORCdivorcedDichotomousVALUEsocial development index by valueContinuousSTRATUMsocioeconomic stratumContinuousQUA_HOUSquality and living spaceContinuousHEALTHACaccess to healthcare and social securityContinuousDULAGeducational lagContinuousDURABdurable goodsContinuous	BMI	body mass index	Continuous
SBPsystolic blood pressureContinuousDBPdiastolic blood pressureContinuousLIV_TOGcommon-law marriageDichotomousMARRIEDmarriedDichotomousSINGLEsingleDichotomousDIVORCdivorcedDichotomousVALUEsocial development index by valueContinuousSTRATUMsocioeconomic stratumContinuousQUA_HOUSquality and living spaceContinuousHEALTHACaccess to healthcare and social securityContinuousDULAGeducational lagContinuous	WC	waist	Continuous
DBPdiastolic blood pressureContinuousLIV_TOGcommon-law marriageDichotomousMARRIEDmarriedDichotomousSINGLEsingleDichotomousDIVORCdivorcedDichotomousVALUEsocial development index by valueContinuousSTRATUMsocioeconomic stratumContinuousQUA_HOUSquality and living spaceContinuousHEALTHACaccess to healthcare and social securityContinuousDULAGeducational lagContinuous	SBP	systolic blood pressure	Continuous
LIV_TOGcommon-law marriageDichotomousMARRIEDmarriedDichotomousSINGLEsingleDichotomousDIVORCdivorcedDichotomousVALUEsocial development index by valueContinuousSTRATUMsocioeconomic stratumContinuousQUA_HOUSquality and living spaceContinuousHEALTHACaccess to healthcare and social securityContinuousDULAGeducational lagContinuous	DBP	diastolic blood pressure	Continuous
MARRIEDmarriedDichotomousSINGLEsingleDichotomousDIVORCdivorcedDichotomousVALUEsocial development index by valueContinuousSTRATUMsocioeconomic stratumContinuousQUA_HOUSquality and living spaceContinuousHEALTHACaccess to healthcare and social securityContinuousDULAGeducational lagContinuous	LIV_TOG	common-law marriage	Dichotomous
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STRATUMsocioeconomic stratumContinuousQUA_HOUSquality and living spaceContinuousHEALTHACaccess to healthcare and social securityContinuousEDULAGeducational lagContinuousDURABdurable goodsContinuous	VALUE	social development index by value	Continuous
QUA_HOUSquality and living spaceContinuousHEALTHACaccess to healthcare and social securityContinuousEDULAGeducational lagContinuousDURABdurable goodsContinuous	STRATUM	socioeconomic stratum	Continuous
HEALTHACaccess to healthcare and social securityContinuousEDULAGeducational lagContinuousDURABdurable goodsContinuous	QUA_HOUS	quality and living space	Continuous
EDULAG educational lag Continuous	HEALTHAC	access to healthcare and social security	Continuous
DURAB durable goods Continuous	EDULAG	educational lag	Continuous
Communication communications	DURAB	durable goods	Continuous
SANITRY sanitary adequacy Continuous	SANITRY	sanitary adequacy	Continuous
ENER_AD energy efficiency Continuous	ENER_AD	energy efficiency	Continuous
ED_LEVEL educational level in the neighborhood Continuous	ED_LEVEL	educational level in the neighborhood	Continuous
SEC_SCHOOL secondary school Dichotomous	SEC_SCHOOL	secondary school	Dichotomous
DOCTORATE doctorate Dichotomous	DOCTORATE	doctorate	Dichotomous
MASTER master Dichotomous	MASTER	master	Dichotomous
SCHOOL school Dichotomous	SCHOOL	school	Dichotomous
BACHELORS bachelor's degree Dichotomous	BACHELORS	bachelor's degree	Dichotomous
HIGH_SCHOOL high school Dichotomous	HIGH_SCHOOL	high school	Dichotomous
TECH_SCHOOL technical school Dichotomous	TECH_SCHOOL	technical school	Dichotomous
NONE no academic degree Dichotomous	NONE	no academic degree	Dichotomous
TOTMET metabolic equivalent of task Continuous	TOTMET	metabolic equivalent of task	Continuous
STAT_ANX state anxiety Dichotomous	STAT_ANX	state anxiety	Dichotomous
TRAIT_ANX trait anxiety Dichotomous	TRAIT_ANX	trait anxiety	Dichotomous
SLPNOTQ sleep was not quiet Continuous	SLPNOTQ	sleep was not quiet	Continuous
BREATH waking up with shortness of breath Continuous	BREATH	waking up with shortness of breath	Continuous
DROWSY feeling drowsy or sleepy Continuous	DROWSY	feeling drowsy or sleepy	Continuous
TROBLS trouble falling asleep Continuous	TROBLS	trouble falling asleep	Continuous
AWAKEN awakens during your sleep time Continuous	AWAKEN	awakens during your sleep time	Continuous
STYAWKE trouble staying awake Continuous	STYAWKE	trouble staying awake	Continuous
TAKENAP takes naps of 5 min or longer Continuous	TAKENAP	takes naps of 5 min or longer	Continuous
SLPD4 sleep disturbances Continuous	SLPD4	sleep disturbances	Continuous
SLPSNR1 snores during sleep Continuous	SLPSNR1	snores during sleep	Continuous
SLPSOB1 sleep short (headache) Continuous	SLPSOB1	sleep short (headache)	Continuous
SLPA2 sleep adequacy Continuous	SLPA2	sleep adequacy	Continuous
SLPS3 somnolence Continuous	SLPS3	somnolence	Continuous
SLPS6 sleep problems (Index I [40]) Continuous	SLPS6	sleep problems (Index I [40])	Continuous
SLPS9 sleep problems (Index II [40]) Continuous	SLPS9	sleep problems (Index II [40])	Continuous
SLPQRAW sleep quantity Continuous	SLPQRAW	sleep quantity	Continuous
SLPOP1 sleep quality Dichotomous	SLPOP1	sleep quality	Dichotomous
SMOKING smoking practice Dichotomous	SMOKING	smoking practice	Dichotomous
CURRENT current smoker Dichotomous	CURRENT	current smoker	Dichotomous

# Table 1. Cont.

Name Variable	Description	Туре
EXSMOKER	ex-smoker	Dichotomous
SMO_PASS	smoker passive	Dichotomous
ALCOHOL	alcohol consumption	Dichotomous
ENERGYDRK	energy drinks	Dichotomous
MOTHEROB	maternal obesity history	Dichotomous
FATHEROB	paternal obesity history	Dichotomous
MOTHERDB	maternal diabetic history	Dichotomous
FATHERDB	paternal diabetic history	Dichotomous
MOTHERHT	maternal hypertension history	Dichotomous
FATHERHT	paternal hypertension history	Dichotomous
MOTHERDL	maternal dyslipidemia history	Dichotomous
FATHERDL	paternal dyslipidemia history	Dichotomous
MOTHERGT	maternal gout history	Dichotomous
FATHERGT	paternal gout history	Dichotomous
URIC	uric acid	Continuous
CREA	creatinine	Continuous
HDLCO	high-density lipoprotein	Continuous
LDLCO	low-density lipoprotein	Continuous
GLU	blood glucose	Continuous
IAT	atherogenic index	Continuous
CHOL_ANT	cholesterol	Continuous
TRIG	triglycerides	Continuous
NA	sodium	Continuous
CALOR	energy	Continuous
PROTEI	total proteins	Continuous
APROT	proteins of animal origin	Continuous
CARBO	carbohydrates	Continuous
SUCR	sucrose	Continuous
FRUCT	fructose	Continuous
LACT	lactose	Continuous
ST	starch	Continuous
MALT	maltose	Continuous
GLU_1	glucose levels based on the dietary survey	Continuous
CRUDE	crude fiber	Continuous
SOLFB	soluble dietary fiber	Continuous
INSFB	insoluble dietary fiber	Continuous
HEMCL	hemicellulose	Continuous
CALC	calcium	Continuous
IKON	total iron	Continuous
MAGN	magnesium	Continuous
ГП V	phosphorus	Continuous
SODIUM	sodium lovels based on the dietary survey	Continuous
7N	zinc	Continuous
CU	copper	Continuous
MN	manganese	Continuous
SE	iodine	Continuous
VITC	vitamin C	Continuous
B1	thiamine	Continuous
B2	riboflavin	Continuous
B6	vitamin B6	Continuous
B12	vitamin B12	Continuous
VITK	vitamin K	Continuous
RETINOL	retinol	Continuous
VITD	vitamin D	Continuous
VITE	vitamin E	Continuous
CHOL_SN	cholesterol levels based on the dietary survey	Continuous
ALCO	alcohol levels based on the dietary survey	Continuous
CAFF	caffeine	Continuous

Name Variable	Description	Туре
AFAT	animal fat	Continuous
VFAT	vegetable fat	Continuous
TFATAV	total fat: animal + vegetable	Continuous
SATFAT	saturated fat	Continuous
MONFAT	monounsaturated fat	Continuous
POLY	polyunsaturated fat	Continuous
MS	MetS	Dichotomous

Table 1. Cont.

### 2.2. Methods

2.2.1. Feature Selection

Feature selection is essential to identify and establish the most critical variables.

In this study, we employed logistic regression to measure the relationship between variables and class alongside machine learning algorithms to discern the most significant features. The algorithms used were RF and RPART (see Machine Learning Modelsbelow), applying the mean decrease accuracy for calculating variable importance, which can be expressed as follows:

$$MDI_{i} = \sum_{all \ nodes} ((Imp(node) - Weight.Imp(node))/NS.N)$$
(1)

where  $MDI_i$  is the mean decrease impurity of the *ith* variable; Imp(node) is the impurity of the node before the split; Weight.Imp(node) is the weighted impurity of the child nodes resulting from the split; and NS.N is the number of samples in the node before the split.

### 2.2.2. Balancing Methods

Balancing methods such as SMOTE and ADASYN have helped address the class imbalance issue within our dataset.

ADASYN (Adaptive Synthetic Sampling), which is part of the UBL R package, takes a unique approach by generating synthetic samples based on the local density of minority class instances, with a focus on instances that are more challenging to learn. In this method, the  $\beta$  parameter controls the desired balance rate between the minority and majority classes during the generation of synthetic samples. When  $\beta$  is set to a value greater than 1, a proportionally larger number of synthetic samples will be generated relative to the instances of the minority class. This further increases the ratio between the minority and majority classes.

The second method, SMOTE (Synthetic Minority Oversampling Technique) of the performanceEstimation R package (Version: 1.1.0), generates synthetic samples for the minority class. In SMOTE, the k parameter determines the number of nearest neighbors used to generate synthetic samples. A small value of k can lead to an excessive generation of synthetic samples that may be too close together, resulting in model overfitting. Moreover, if k is too large, synthetic samples may be less representative of the minority class and fail to capture data variability adequately.

# 2.2.3. Machine Learning Models

To build the models, we applied two machine learning algorithms, RF [41,42] and RPART [43,44], as well as PCA [45,46]. RF, introduced by Breiman [47], is a machine learning algorithm combining multiple decision trees to create a model with the highest accuracy. Rpart (Recursive Partitioning and Regression Trees), by Breiman [48], works by recursively partitioning the input data based on predictor variables to create a tree-like structure. This algorithm aims to find the optimal splits in the data that maximize the homogeneity or purity of the resulting subgroups. Principal component analysis (PCA) is a data analysis technique used to simplify the complexity of data by reducing their dimensionality, facilitating visualization and analysis.

### 2.3. Performance Measures

We used sensitivity, specificity, and balanced accuracy (B.ACC) to evaluate model performance. These metrics provide a fair assessment of the model's performance across all classes, considering the issue of class imbalance.

$$SENS = \frac{TP}{TP + FN}$$
(2)

$$SPC = \frac{TN}{FP + TN} \tag{3}$$

$$B.ACC = \left(\frac{1}{2}\right) \left(\frac{TP}{P} + \frac{TN}{N}\right) \tag{4}$$

where *P* = *Positive*, *N* = *Negative*, *TP* = *True Positive*, *FN* = *False Negative*, *TN* = *True Negative*, *and FP* = *False Positive*, respectively.

# 3. Statistical Analysis and Development of Prediction Models

All experiments were performed using the R programming language (3.6.1) [49]. Min-max was used to normalize continuous variables, and dichotomous variables were represented as numbers. Figure 1 provides a general overview of the experimental process described in this section. To develop predictive models, it was necessary to process the data and implement a balancing technique. The minority class was oversampled, taking into account the majority class. As a first step, SMOTE was applied, and it was necessary to determine the best value of *k* (number of nearest neighbors), so experiments were conducted by varying *k* (here, we present k = 1, k = 5, and k = 9). In this process, the dataset was randomly divided into 70% for training and 30% for testing. To accomplish this task, we applied two machine learning algorithms, RF and RPART. In the case of RF, we varied the *mtry* parameter from 1 to 10 and considered *ntree* values of 100, 300, 500, and 1000 for each model.

Additionally, a subset of features was extracted in each created model using the variable importance (VarImp) of RF, and a 10-fold cross-validation was performed. Similarly, in the case of RPART, parameter tuning was conducted by considering cp = 0, cp = 0.05, and cp = 0.005, using a 10-fold cross-validation. Likewise, a subset of features was extracted in each created model.

Once the feature subsets were obtained, along with the optimal value for each corresponding parameter of each algorithm and data balancing technique, we tested the generated feature subsets using RF and RPART. This was accomplished by conducting 30 runs with different seeds to assess the performance of each model. In all experiments, a minimum of 30 independent runs were conducted for each algorithm using 30 different seeds. The mean and standard deviation of the performance measures were calculated for each of these runs.



Figure 1. Experimental process.

# 4. Results

Understanding how MetS, nutrition, sleep disturbances, and SDI relate in men and women can have important clinical and public health implications. In this study, we used logistic regression before dataset balancing to pinpoint the critical variables associated with MetS in both sexes. Table 2 presents the results of the features and their corresponding values obtained.

	Women			Men			
Variable	Coefficient	<i>p</i> _Value	Variable	Coefficient	<i>p</i> _Value		
GLU	4.61438598	$6.24  imes 10^{-59}$	GLU	3.94711748	$2.45  imes 10^{-39}$		
TRIG	3.63418178	$1.18  imes 10^{-37}$	TRIG	2.98165065	$3.25  imes 10^{-24}$		
WC	1.75532078	$2.86 \times 10^{-9}$	WC	2.53131848	$1.02  imes 10^{-9}$		
BMI	1.60919304	$1.05  imes 10^{-6}$	IAT	2.06238741	$5.13  imes 10^{-11}$		
SBP	1.40299133	$1.15  imes 10^{-12}$	SBP	1.53063308	$1.31  imes 10^{-11}$		
PROTEI	0.90748897	0.08529715	B12	1.41903991	0.00880359		
FRUCT	0.73077934	0.23874313	BMI	1.40229014	0.00087404		
CHOL_SN	0.72037259	0.06868106	LACT	1.29691863	0.00581383		
URIC	0.65547784	0.01333401	CARBO	1.18935354	0.0886463		
CU	0.64813271	0.17111299	GLU_1	1.1674073	0.10024746		

Table 2. Features and values obtained through logistic regression for men and women.

Analyzing the data, in men, the top 10 variables most related to MetS are GLU, TRIG, WC, IAT, SBP, vitamin B12 (B12), BMI, lactose (LACT), carbohydrates (CARBO), and high glucose levels based on the dietary survey (GLU\_1). Conversely, in women, the ten most relevant variables include GLU, TRIG, WC, BMI, SBP, total proteins (PROTEI), fructose (FRUCT), high cholesterol total based on the dietary survey (CHOL\_SN), URIC, and copper (CU). To achieve a more effective visualization of these prominent features from the logistic regression for both men and women, Figure 2 is presented. Red square symbols represent the most substantial variables for women, while blue triangles represent those for men. A cautionary note must be made for the seemingly outlier behavior of blood glucose and triglycerides with very high coefficients. Let us recall that these features are closely related to the very definition of MetS. Such variables were included in our models only for the sake of database completeness and comprehensiveness. Detailed results for women can be found in Supplementary Table S1, and those for men are available in Supplementary Table S2.



**Figure 2.** The most important variables obtained through logistic regression for men and women before data balancing.

Subsequently, we employed SMOTE and ADASYN with RF and RPART to reassess the most influential features associated with MetS prediction within a now balanced dataset. Following this, with the data balancing techniques effectively applied and their parameters fine-tuned, we extract feature subsets by utilizing RPART and RF for both women and men.

Extracting features related to MetS in a balanced dataset improves model generalization (conducting training more evenly and accurately), optimizing performance, and reducing overfitting. Considering the challenges associated with including all variables in a model, such as noise, redundancy, and overfitting, we extracted the 17 variables with the highest values obtained in each model of RF and RPART after applying SMOTE and ADASYN.

The extracted feature subsets, along with their respective values, are presented in Tables 3-6. These tables also detail the employed balancing technique for each set of variables and their corresponding parameters ranging from 1 to 5. Each subset was adjusted for its corresponding parameter, B for ADASYN and *k* for SMOTE, considering values of 1 and 5.

Similarly, Table 7 showcases the performance achieved by the RF algorithm, while Table 8 presents the performance of the RPART algorithm. In both tables, the Value column provides information regarding the relative importance of each feature.

Table 3. Features of men obtained using RF with ADASYN and SMOTE applied.

ADASYN, $B = 1$		ADASYN, $B = 5$		SMOTE, K = 1	SMOTE, K = 1		SMOTE, K = 5	
Features	Value	Features	Value	Features	Value	Features	Value	
BMI	92.9499	ENER_AD	130.906694	MOTHERDL	204.657628	BMI	289.868211	
WEIGHT	49.4782	BMI	104.213511	ALCOHOL	199.602686	MOTHERDL	172.071267	
ENER_AD	48.8887	WEIGHT	81.5087781	BMI	198.579371	WEIGHT	169.929592	
EDULAG	45.2797	EDULAG	67.7406035	SLPSOB1	111.323472	ALCOHOL	131.283664	
LIV_TOG	33.3601	ALCOHOL	62.4379604	CURRENT	95.3509822	IAT	93.2909179	
DURAB	31.5583	STRATUM	57.134903	BREATH	80.8262246	CHOL_ANT	63.4703128	
MOTHERGT	27.5583	ED_LEVEL	55.578244	SLPD4	70.1756789	NA	49.2933568	
IAT	25.7470	NONE	38.1101529	CAFF	68.9892898	CREA	45.8846962	
HEALTHAC	23.4522	DURAB	36.4129389	SLP6	60.2949079	SINGLE	44.6897663	
DIVORC	20.1163	VALUE	36.0130176	WEIGHT	56.9297661	SLPSNR1	35.672622	
QUA_HOUS	17.4925	DIVORC	35.8243538	TOTMET	52.4806201	MOTHERDB	35.21356	
STRATUM	16.1269	FATHERGT	33.7033121	ALCO	45.7609412	ENERGYDRK	34.0359073	
FATHERGT	14.5872	MASTER	29.8751736	AWAKEN	39.0795326	URIC	31.8268793	
NONE	14.0213	PRIMARIA	28.3852397	IAT	38.042823	AGE	27.9839119	
MARRIED	13.9584	SLPSNR1	27.9671847	TROBLS	36.7528999	MARRIED	27.8864259	
VALUE	13.8059	AGE	24.3706018	STYAWKE	36.2387269	DOCTORATE	24.4733499	
URIC	13.7930	IAT	22.0506592	MALT	34.3472852	DIVORC	24.142464	
SANITRY	13.5609	SANITRY	21.924077	BACHELORS	33.7934562	SLPOP1	23.8868609	
SINGLE	13.4148	SINGLE	21.7818986	MARRIED	32.6228111	SEC_SCHOOL	22.755325	
ALCOHOL	12.9798	DOCTORATE	19.8069099	SLP9	31.0845509	SLPQRAW	20.666244	

Table 4. Features of men obtained using RPART with ADASYN and SMOTE applied.

ADASYN, B = 1 ADASYN, B = 5		: 5	SMOTE, K = 1	SMOTE, K = 1		SMOTE, K = 5	
Features	Value	Features	Value	Features	Value	Features	Value
LIV_TOG	447.069761	BMI	683.735277	BMI	185.940586	BMI	164.086828
BMI	402.975487	ENER_AD	619.998675	WEIGHT	131.361866	WEIGHT	132.276557
ENER_AD	338.664389	EDULAG	565.325738	FATHERGT	115.496204	IAT	131.937059
EDULAG	325.498647	ALCOHOL	355.970533	MOTHERDL	96.1708037	SINGLE	83.6531675
DURAB	285.861702	WEIGHT	295.254303	IAT	67.2839991	MOTHERDL	71.6947353
SLP6	64.2112969	DIVORC	214.489844	AGE	40.9532174	APROT	47.2274885
WEIGHT	33.1175418	NONE	200.599299	LACT	28.7681412	TFATAV	22.4867652
IAT	27.5407406	MOTHERGT	178.450647	MOTHERHT	25.3414479	ST	20.7519258
FATHEROB	14.5734264			PROTEI	14.5865884	HEALTHAC	19.7752349
SLPSNR1	13.7361635			CAFF	14.1658755	SATFAT	17.5962564
				ZN	12.4515539	HEIGHT	16.3718359
				MN	12.20696	CHOL_ANT	15.4222905
				IRON	10.5317678	MONFAT	13.9908309
				VALUE	10.2017285	CREA	13.6358167
				STYAWKE	10.1887194	URIC	11.0085972
				MONFAT	10.0410598	AGE	10.5421496
				CHOL_ANT	9.78675973	CALC	10.0034374
				ST	9.41791645	SMOKING	9.53883547
				SINGLE	9.40405705	LACT	9.34161011
				SOLFB	7.74765092	TOTMET	9.09355989

ADASYN, $B = 1$		ADASYN, $B = 5$		SMOTE, K = 1		SMOTE, K = 5	
Features	Value	Features	Value	Features	Value	Features	Value
BMI	208.269603	ENER_AD	344.249674	WEIGHT	321.316267	BMI	484.307061
IAT	151.849516	BMI	210.90055	IAT	294.958989	IAT	481.475021
WEIGHT	98.3094923	IAT	173.895403	BMI	253.281611	WEIGHT	339.174822
EDULAG	98.0933243	ALCOHOL	146.230976	EXSMOKER	246.78181	URIC	142.754087
LIV_TOG	82.4204188	DURAB	142.91494	MASTER	241.332636	SLPSNR1	92.0496746
ENER_AD	80.7154997	EDULAG	142.817907	FATHERDL	211.443455	CHOL_ANT	74.3706077
URIC	60.4722703	WEIGHT	128.038926	CREA	170.195583	AGE	72.769531
VALUE	53.5122927	VALUE	80.989846	MOTHERHT	125.867318	SLPSOB1	70.1959444
DURAB	48.2486067	NONE	76.4699068	SLPSOB1	125.384246	BREATH	60.3028803
QUA_HOUS	37.8080123	QUA_HOUS	62.8303545	SMO_PASS	86.2763209	TRAIT_ANX	56.4099594
SLPSNR1	31.399627	BACHELORS	56.0706757	BREATH	83.1176663	SMO_PASS	50.8288614
HEALTHAC	30.6724986	SANITRY	52.5802813	CHOL_ANT	78.8668934	SANITRY	50.3648334
SANITRY	24.2597947	HEALTHAC	45.9188536	SMOKING	57.7946015	MOTHERDL	50.0567677
ALCOHOL	24.2064626	URIC	43.8531276	TRAIT_ANX	57.3909833	DROWSY	44.564559
AGE	21.594859	SINGLE	39.5694722	SLPSNR1	51.1574483	SMOKING	44.5264858
SINGLE	18.0193809	DIVORC	37.3860944	NA	50.3156936	SINGLE	41.993735
HIGH_SCHOOL	17.1684616	AGE	33.8392029	MARRIED	48.4664641	EXSMOKER	38.9120379
SLP6	16.0530682	TECH_SCHOOL	32.4092154	SLPOP1	48.3006717	SEC_SCHOOL	38.4719692
SOLFB	14.4271683	SCHOOL	28.2955057			SLPNOTQ	35.6761924
FATHERGT	13.8839264	MARRIED	27.6425229			~	

 Table 5. Features of women obtained using RF with ADASYN and SMOTE applied.

Table 6. Features of women obtained using RPART with ADASYN and SMOTE applied.

ADASYN, B =	1	ADASYN, $B = 5$		SMOTE, K = 1		SMOTE, K = 5	
Features	Value	Features	Value	Features	Value	Features	Value
BMI	664.323812	BMI	1164.1686	BMI	427.45413	IAT	483.233069
LIV_TOG	535.392713	DURAB	1117.88127	IAT	363.893488	BMI	410.367827
ENER_AD	507.53479	ENER_AD	1090.27197	SLPSNR1	259.475806	WEIGHT	409.777127
EDULAG	505.45874	EDULAG	772.049538	SLPS3	259.475806	URIC	278.65513
IAT	468.310602	ALCOHOL	655.016952	EXSMOKER	217.54026	SLPSNR1	86.0218576
		NONE	533.217568			SMOKING	31.3976405
		IAT	380.443927			SLPS3	30.5201011
		WEIGHT	366.577281			SODIUM	15.7251124
		VALUE	104.231729			ALCOHOL	12.4735987
		TECH_SCHOOL	92.1094015			SATFAT	12.1523683
						MONFAT	12.1446951
						NA	11.2712045
						VITE	10.3455105
						CHOL_ANT	9.04441276
						FATHERDB	8.09870623
						SUCR	7.16739885
						MARRIED	6.39473684
						FRUCT	4.94398493
						MALT	4.8372105

# Table 7. Results of the random forest models applying ADASYN and SMOTE in men and women.

Sex	Subset	Parameters	B.ACC (%)	Sensitivity (%)	Specificity (%)
Men	ADASYN, $B = 1$	Mtry = 9	86.22	90.93	81.50
		Ntree $= 200$	$\pm 0.26$	$\pm 0.60$	$\pm 0.41$
Men	ADASYN, $B = 5$	Mtry = 8	85.56	87.85	83.26
		Ntree = 200	$\pm 0.34$	$\pm 0.49$	$\pm 0.55$
Men	SMOTE, $K = 1$	Mtry = 10	82.86	91.51	74.21
		Ntree $= 200$	$\pm$ 1.66	$\pm 0.68$	$\pm$ 3.45
Men	SMOTE, $K = 5$	Mtry = 10	75.43	90.48	60.39
		Ntree $= 100$	$\pm$ 1.29	$\pm 0.95$	$\pm 2.50$
Women	ADASYN, $B = 1$	Mtry = 10	87.12	91.10	83.15
		Ntree $= 200$	$\pm 0.25$	$\pm 0.40$	$\pm 0.29$
Women	ADASYN, $B = 5$	Mtry = 10	86.73	88.62	84.84
		Ntree = 300	$\pm 0.20$	$\pm 0.24$	$\pm 0.36$
Women	SMOTE, $K = 1$	Mtry = 10	82.55	90.48	74.62
		Ntree $= 300$	$\pm 0.71$	$\pm 0.39$	$\pm 1.46$
Women	SMOTE, $K = 5$	Mtry = 10	88.50	91.91	85.10
		Ntree $= 300$	$\pm 0.40$	$\pm 0.42$	$\pm 0.75$

Sex	Subset	Parameters	B.ACC (%)	Sensitivity (%)	Specificity (%)
Men	ADASYN, $B = 1$	cp = 0.05	82.14	81.57	82.71
		•	$\pm 1.75$	$\pm$ 3.38	$\pm$ 2.07
Men	ADASYN, $B = 5$	cp = 0.05	82.32	82.87	81.77
			$\pm 0.99$	$\pm$ 4.67	$\pm$ 5.02
Men	SMOTE, $K = 1$	cp = 0.001	75.41	73.09	77.73
			$\pm 2.78$	$\pm$ 4.07	$\pm$ 5.36
Men	SMOTE, $K = 5$	cp = 0.002	74.67	71.96	77.38
			$\pm 2.78$	$\pm$ 4.07	$\pm$ 5.36
Women	ADASYN, $B = 1$	cp = 0.05	78.90	69.96	87.84
			$\pm 0.31$	$\pm 0.00$	$\pm 0.62$
Women	ADASYN, $B = 5$	cp = 0.05	78.90	69.96	87.84
			$\pm 0.31$	$\pm 0.00$	$\pm 0.62$
Women	SMOTE, $K = 1$	cp = 0.001	80.86	79.85	81.87
			$\pm$ 1.91	$\pm$ 3.79	$\pm$ 3.57
Women	SMOTE, $K = 5$	cp = 0.005	84.49	84.20	84.79
			$\pm 1.43$	$\pm$ 3.01	$\pm 2.51$

Table 8. Results of the RPART models applying ADASYN and SMOTE in men and women.

### 4.1. Best Features for Men Using RF and ADASYN/SMOTE

Specifically, Table 3 exhibits four feature subsets obtained from male data using random forest with ADASYN and SMOTE. According to Table 7, the most effective subset was obtained by applying ADASYN with B = 1 with a balanced accuracy of 86.22% and a deviation standard of 0.26%.

The most influential factor within this subset was BMI, which had a significant importance value of 92.9499. This was followed by WEIGHT and energy efficiency (ENER\_AD), with importance values of 49.4782 and 48.8887, respectively. Other factors such as educational lag (EDULAG), common-law marriage (LIV\_TOG), durable goods (DURAB), and maternal gout history (MOTHERGT) also contributed to the model, albeit to a lesser extent.

### 4.2. Best Features for Men Using RPART and ADASYN/SMOTE

In the case of features obtained by RPART (see Table 4), using both SMOTE and ADASYN, the results were slightly worse than those obtained with RF (Table 3). In this scenario, the best subset was achieved by the subset with the parameter ADASYN = 5, which achieved an 82.32% balanced accuracy metric with a standard deviation of 0.99% (see Table 8).

Switching gears to the outcomes yielded by random forest with ADASYN using a B value of 5, BMI takes center stage with a substantial value of 683.74, signifying its paramount role in predicting the outcomes related to the examined condition. Following closely in significance are ENERGY\_AD and EDULAG, boasting values of 619.99 and 565.33, respectively, both making substantial contributions to predictive capability. ALCOHOL and WEIGHT also exhibit noteworthy importance with values of 355.97 and 295.25, underlining their relevance within the model. Moreover, features like divorce (DIVORC), no academic degree (NONE), and MOTHERGT, while exerting a comparatively lower influence, still contribute to the model's predictive capacity, as indicated by their respective values.

#### 4.3. Best Features for Women Using RF and ADASYN/SMOTE

The random forest model using SMOTE with k = 5 achieved the best performance for women, reaching an 88.50% accuracy with a standard deviation of 0.40% (see Table 7). In this case, Table 5 reveals that BMI was identified as the primary predictor, with a notable value of 484.31, clearly highlighting the critical importance of BMI in predicting MetS in this particular context. Additionally, IAT (481.48) and WEIGHT (339.17) also showed significant associations, further emphasizing the relevance of weight-related measurements.

Including sleep disturbances (SLPSNR1, SLPSOB1, BREATH, DROWSY, and SLP-NOTQ) and even cholesterol levels (CHOL\_ANT) among the influential variables underscores their pivotal contributions to MetS prediction in women. The importance of AGE and SDI parameters like sanitary adequacy (SANITRY) is also noteworthy. It is essential to highlight that psychological factors such as trait anxiety (TRAIT\_ANX) were included, accounting for the potential influence of mental health aspects in MetS prediction.

### 4.4. Best Features for Women Using RPART and ADASYN/SMOTE

In this instance, SMOTE with k = 5, combined with RPART, achieved the best performance, attaining a balanced accuracy of 84.49% with a standard deviation of 1.43% (see Table 8). The results of the corresponding subset (RPART applied to women's data using SMOTE with a parameter value, k = 5) shown in Table 6 reveal that the most influential feature was IAT, with a value of 483.23, followed closely by BMI and WEIGHT, which have values of 410.37 and 409.78, respectively. Features like URIC, snores during sleep (SLPSNR1), somnolence (SLPS3), SODIUM, vitamin E consumption (VITE), and habitual smoking (SMOKING) also exhibit noticeable influence, indicating their relevance in understanding the targeted phenomenon. Conversely, some nutrients like sucrose (SUCR), maltose (MALT), and FRUCT have relatively lower values; however, they can provide valuable information about dietary habits, nutritional deficiencies, or behaviors related to MetS.

This study's results, employing random forest and RPART algorithms and SMOTE and ADASYN techniques for both genders, offer valuable insights. These results underscore the importance of health and lifestyle elements in MetS prediction, encompassing sleep disturbances, cholesterol levels, age, psychological factors, and SDI parameters.

# 4.5. Analyzing the Best Features Using PCA

Based on the results of the features obtained in the best models, we used PCA to visually and graphically analyze the top features for men and women to explore potential correlations and latent patterns among these influential factors and reduce dimensionality to the greatest possible extent.

In the case of men, we considered feature subsets obtained from the random forest model using ADASYN with B = 1 and RPART with ADASYN and B = 5. The subsequent features were integrated: BMI, WEIGHT, ENER\_AD, EDULAG, LIV\_TOG, DURAB, MOTHERGT, IAT, HEALTHAC, DIVORC, QUA\_HOUS, STRATUM, FATHERGT, NONE, MARRIED, VALUE, URIC, SANITRY, SINGLE, and ALCOHOL.

For women, we considered feature subsets obtained from the random forest model with SMOTE and k = 5 and the RPART model with SMOTE and k = 5. These models are regarded because they achieved the highest performance (see Tables 7 and 8 where extremely small percentage uncertainty values in Table 8 are shown rounded down to 0.00 for clearer presentation). The following features were included: BMI, IAT, WEIGHT, URIC, SLPSNR1, CHOL\_ANT, AGE, SLPSOB1, BREATH, TRAIT\_ANX, SMO\_PASS, SANITRY, MOTHERDL, DROWSY, SMOKING, SINGLE, EXSMOKER, SEC\_SCHOOL, SLPNOTQ, SLPS3, SODIUM, ALCOHOL, SATFAT, MONFAT, NA, VITE, FATHERDB, SUCR, MARRIED, FRUCT, ZN, and MALT.

The PCA analysis, as shown in Figure 3, revealed the relative importance of features concerning MetS in men. The first principal component (PC1) was more influenced by features such as WEIGHT, BMI, and SDI by value (VALUE), suggesting that these variables significantly contributed to the observed variability in the data. On the other hand, the second principal component (PC2) was more affected by features like EDULAG and socioeconomic stratum (STRATUM). These findings indicated that weight and BMI were prominent factors in the context of MetS, as well as education and socioeconomic stratum. In this case, PC1 was considered the most significant component, as it had a magnitude of 0.508501, capturing most of the variability, while PC2 had a magnitude of 0.499809.



Figure 3. PCA of features of men for metabolic syndrome with clusters.

On the other hand, in the case of women (see Figure 4), features associated with the variability of MetS along PC1 were sodium levels based on the dietary survey (SODIUM), saturated fat (SATFAT), and monosaturated fat (MONFAT), which exhibit significant magnitudes in PC1. Furthermore, BMI significantly influences PC1, indicating its association with this variability. Conversely, variables like short sleep duration (SLPSOB1) and waking up with shortness of breath (BREATH) demonstrate significant magnitudes in PC2. Similarly, TRAIT\_ANX and feeling drowsy or sleepy (DROWSY) are associated with variability in PC2. Therefore, considering the magnitudes in the principal components, the features in women associated with the risk of MetS include SODIUM, SATFAT, and MONFAT from PC1, as well as SLPNOTQ and SLPSOB1 from PC2.



Figure 4. PCA of features of women for metabolic syndrome with clusters.

# 5. Discussion

MetS is a severe and potentially life-threatening condition that significantly increases the risk of developing cardiovascular diseases and also increases the severity of diabetes. Over the years, several consistently highlighted risk factors have been associated with MetS. Considering imbalanced data, this study analyzed participant data from a cohort to identify the primary risk factors in both men and women. Subsequently, data balancing techniques were applied to ascertain whether significant differences exist, contributing to selecting risk factors for MetS prediction. Using data balancing techniques is crucial in this context, as it helps ensure a more accurate and unbiased identification of relevant risk factors, especially when working with unevenly distributed data. In this study, we applied logistic regression to identify the risk factors in men and women that predict the occurrence of MetS within an imbalanced data environment.

### 5.1. Logistic Regression

The logistic regression analysis in women demonstrates (as expected, of course) the strong connection between MetS and elevated glucose levels, which is in line with prior research [50,51], emphasizing the crucial role of glucose in MetS. Additionally, uric acid is also identified as a significant risk factor in women [52–54]. Subsequent findings revealed other risk factors, including waist circumference, BMI, and systolic blood pressure, which are all essential components of MetS. WC is an indicator of abdominal obesity closely linked to insulin resistance, while BMI reflects the relationship between weight and height, which is a significant obesity-related risk factor for MetS.

Furthermore, Figure 2 highlights additional significant factors derived from dietary data, including the intake of protein and fructose [55–57]. When these two nutrients are combined, they have been linked to an elevated risk of MetS [58]. Likewise, copper consumption is evident, which can impact glucose regulation [3] and liver function, which

are both crucial components in MetS [59]. These factors underscore the importance of moderate consumption of these nutrients in preventing MetS.

In the case of men, glucose was identified as the primary factor associated with MetS, followed by triglycerides, waist circumference, atherogenic index, and systolic blood pressure. Additionally, the consumption of lactose [60] and carbohydrates [61] was noted among the nutrients. Elevated glucose, triglycerides, and waist circumference are critical markers of MetS, while the atherogenic index assesses cardiovascular risk. High systolic blood pressure is another significant component of this syndrome. Regarding lactose, it is worth noting that certain dairy products may include added sugars, which can potentially increase the overall calorie intake [62]. This potentially contributes to obesity and insulin resistance, which are two critical factors in the onset of MetS. Moreover, high lactose consumption is associated with a risk factor for developing diabetes, cardiovascular diseases, and increased cholesterol levels [63,64].

It is possible that when working with unbalanced datasets, machine learning models like logistic regression tend to be biased towards the majority class. For this reason, data balancing techniques such as SMOTE and ADASYN were used to enable a more equitable training of the models to identify more precise relationships between variables and the MetS.

# 5.2. Use of Machine Learning with Synthetic Data

The most effective machine learning models for women revealed associations with attributes related to sleep quality, such as snores during sleep [65], short sleep duration (SLPSOB1) [66], waking up with shortness of breath (BREATH) [67], restless sleep (SLP-NOTQ) [68], and somnolence (SLPS3). Multiple studies have shown that poor sleep quality is closely linked to cardiovascular disease [69,70], diabetes [71], and MetS [72], as well as other adverse health outcomes. In the case of women, an increased likelihood of facing significant risks related to cardiovascular diseases and sleep problems has been observed, especially for those in the postmenopausal stage, which, in turn, can contribute to the development of risks associated with MetS [73]. Additionally, they highlighted factors related to anxiety (TRAIT\_ANX), despite the association between MetS and anxiety remaining a subject of debate due to various issues [74], this study, like some others [75–78], identified anxiety as one of the critical factors that predisposing women to MetS.

In the same way, ex-smokers and current smokers (EXSMOKER, SMOKING) were found to be relevant features; based on this, it has been observed that both smokers and former smokers are predisposed to MetS. This finding is supported by various studies that suggest that smoking can have an adverse impact on blood lipid levels and lead to metabolic disturbances [79–81].

In women, nutritional components also appeared as relevant features, such as SATFAT, MONFAT, SUCR, FRUCT, and MALT. Based on this, a study has revealed that fructose, sucrose, and maltose are critical components of the leading nutrient pattern associated with a higher risk of MetS [58].

In the case of men, the most effective machine learning models displayed more pronounced associations with features linked to the SDI, encompassing ENER\_AD, EDULAG, durable goods (DURAB, HEALTHHAC), quality and living space (QUA\_HOUS), socioeconomic stratum (STRATUM), social development index by value (VALUE), and sanitary adequacy (SANITRY). In studies [22,82–84], a significant association has been observed between a low socioeconomic level and the prevalence of metabolic syndrome. Furthermore, these models underscored variables related to parental gout conditions (MOTHERGT, FATHERGT). This supports research exploring the genetic predisposition to gout and suggests that a family history of this disease may increase the risk of other family members developing it [85]. This condition may also be related to metabolic syndrome due to poor dietary habits that could lead to obesity and insulin resistance [86,87].

### 5.3. Principal Component Analysis

Based on the resulting features obtained for men and women via machine learning models, we applied principal component analysis to identify trends and potential correlations. The PCA conducted using the features obtained for men (Figure 3) showed that *PC1* (the most significant component) revealed a strong association of body-related factors, specifically WEIGHT, and BMI. *PC2* shows a strong correlation among variables related to the SDI. This indicates that the SDI plays a significant role in the onset of MetS, in addition to focusing on interventions related to weight and obesity management.

Figure 3 depicts the distribution of participants in clusters, where Cluster 1, highlighted in green, turned out to be the cluster most predisposed to developing MetS. The arrows emphasize the contribution of individual features to the principal components.

In the context of MetS in women, the most influential factors in *PC1* were factors related to dietary components such as sodium levels based on the dietary survey (SODIUM), SATFAT, and monounsaturated fats (MONFAT), sucrose (SUCR), and FRUCT, among others. *PC2* exhibits a trend towards variables related to poor quality of sleep and anxiety, as SLPSOB1, TRAIT\_ANX, SLPNOTQ, and SLPS3 have significant values in this component. Other variables related to smoking and education (SEC\_SCHOOL) also notably influence this component. This suggests that dietary control is crucial in preventing MetS among women, as well as addressing poor sleep quality and anxiety. Hence, PCA highlights relevant differences in the presentation and risk factors of MetS between men and women [88,89], which is an issue that is progressively gaining relevance in the biomedical literature [90].

The PCA results for women illustrated in Figure 4 show the distribution of participants in clusters. Similarly to the men's analysis, the cluster most predisposed to developing MetS was Cluster 1, which is depicted by yellow dots.

### 5.4. Implications for Metabolic Syndrome Surveillance, Risk Factors, and Public Health Policy

The results of this project suggest several key findings related to the diagnosis of metabolic syndrome:

- Identification of known risk factors: For both men and women, specific variables were identified as strongly related to MetS. These included glucose (GLU), triglycerides (TRIG), waist circumference (WC), body mass index (BMI), and systolic blood pressure (SBP), among others. Notably, these variables are consistent with established criteria for diagnosing MetS, reflecting their importance in understanding the condition.
- 2. Gender-Specific Influential Factors: This study highlights that certain factors vary in importance between men and women in predicting MetS. For instance, vitamin B12, lactose, and carbohydrates were influential in men, while total proteins, fructose, and copper were significant for women. These gender-specific variations underscore the complexity of MetS and the need for tailored diagnostic approaches. One cautionary note regarding potential outliers, specifically blood glucose and triglycerides, emphasizes their close association with the definition of MetS.
- 3. Influence of Sleep and Dietary Habits: The inclusion of sleep-related variables (sleep disturbances, breathing issues) and dietary elements (cholesterol levels, nutrients) underscores their relevance in predicting MetS. These findings suggest that lifestyle factors and dietary habits are integral components in the diagnostic considerations for MetS.
- 4. Potential Role of Psychological Factors: Psychological factors such as trait anxiety were included in the analysis, emphasizing the potential influence of mental health aspects in predicting MetS for both men and women.
- Gender-Specific Dietary Influences: For women, the analysis identified specific dietary factors like sodium levels, saturated fat, and monounsaturated fat as influential. This emphasizes the importance of considering gender-specific dietary influences in MetS diagnosis.

Understanding the gender-specific variations and influential factors highlighted in this study can inform targeted interventions that address the unique needs of both men and women. Public health policies can be crafted to recognize and address the gender-specific variations in the risk factors for MetS. By tailoring interventions to the specific needs of each gender, policymakers can enhance the effectiveness of preventive measures. Moreover, this study underscores the importance of lifestyle factors, including sleep patterns and dietary habits, in predicting MetS. Public health initiatives can thus prioritize educational campaigns and interventions promoting healthier sleep practices and balanced diets. Encouraging regular physical activity, reducing sedentary behaviors, and emphasizing the significance of maintaining a healthy weight can be integral components of public health programs aimed at preventing MetS.

Given the inclusion of psychological factors such as trait anxiety in the analysis, public health policies can integrate mental health considerations into MetS prevention strategies. Mental health awareness campaigns, stress management programs, and access to mental health resources can contribute to holistic approaches addressing the interconnectedness of mental and physical well-being. Public health campaigns can leverage the study's findings to engage communities and raise awareness about the risk factors associated with MetS. Community-based initiatives can offer educational resources, workshops, and screenings to empower individuals to make informed lifestyle choices. By fostering a culture of health consciousness and providing accessible information, public health policies can contribute to the early detection and prevention of MetS. In view of the evolving nature of health trends and behaviors, public health policies should include mechanisms for continuous monitoring and adaptation. Regular assessments of the population's health status, behavior patterns, and response to interventions can inform policy adjustments. This dynamic approach ensures that public health strategies remain effective and responsive to changing circumstances.

#### 5.5. The Role of Social Development Dimensions in Metabolic Syndrome

It is worthwhile to recall that after applying balancing techniques, relevant associations arise between metabolic syndrome and some SDI dimensions (see Figure 5). These effects are moderate-to-medium-sized yet statistically significant. Indeed, since some of these aspects may be modifiable by public policy, it is relevant to consider them. Metabolic syndrome has been previously reported to be related to social dimensions and inequality, but also to dietary patterns [91,92]. Interestingly Soofu and coworkers [91] also report the effect that we found of an association of MetS to housing conditions and ownership of durable assets. Inadequate housing conditions, in particular, have been discussed to contribute to an increase in the risk of cardiovascular disease [93]. In fact, local residential environments may constitute significant risk factors for MetS, which is a fact that needs to be considered in order to develop environmental interventions to improve population health [94].

Restricted access to education (referred to as EDULAG in Figure 5) has also been considered a relevant feature related to MetS [95,96]. Indeed, education levels have been found to be among the best predictors of metabolic conditions in another Mexico City cohort [97]. A similar association has been reported with regards to housing (QUA\_HOUS in Figure 5) [98,99]. A study in an urban Korean population found that non-apartment residents were more likely to have MetS and related phenotypes compared to apartment residents in a model that was adjusted for confounding variables such as sociodemographic characteristics, residence area, health behavior, and nutritional information awareness [93]. Sanitary conditions are known to modify both environmental conditions and even intrinsic factors such as the gut microbiota, affecting the development of MetS [100–102]. All of these dimensions of social development are related in a non-trivial fashion to the development of the complex pathophenotypes making up metabolic syndrome as is further evidenced by our study. However, the actual relationships between these and other risk factors remain to



be investigated as open questions that must be studied in order to design targeted public health interventions.

**Figure 5.** Top features for men and women considering the results of RF and RPART applying balancing techniques.

# 6. Conclusions

In this study, logistic regression was initially utilized to identify pivotal factors linked to MetS across genders, followed by dataset balancing techniques. Our findings indicated significant variables for men, including high glucose levels, triglycerides, waist circumference, systolic blood pressure, vitamin B12, body mass index, high intake of carbohydrates, and lactose. For women, critical factors were glucose levels, triglycerides, waist circumference, body mass index, systolic blood pressure, total protein intake, fructose, cholesterol, uric acid, and copper levels. Further analysis employing SMOTE and ADASYN with RF and RPART methods re-evaluated critical features for MetS prediction in a balanced dataset. This improved model generalization by ensuring more consistent and precise training, enhancing performance, and minimizing overfitting risks. Notably, the analysis also highlighted the relevance of family history of gout as a significant factor, particularly among men. This finding underscores the potential genetic predisposition to gout, suggesting that a familial history of the condition might increase the likelihood of MetS in relatives, possibly due to shared dietary habits contributing to obesity and insulin resistance. These insights emphasize the need for gender-specific public health strategies and medical interventions, considering both the common risk factors and those unique to each gender, such as the family history of gout, to effectively manage and prevent MetS.

# Limitations

The current study has some limitations. This research was based on data from a cohort of relatively healthy adult residents of Mexico City. The regional emphasis of the study might affect generalizability; therefore, it is advisable to exercise caution when extrapolating the findings to wider populations. All data on socioeconomic status, lifestyle habits, family medical history, and macro- and micronutrient intake were self-reported. Although we trust the veracity of the information, some details may have been omitted or not remembered by the participants. Likewise, the instruments applied to evaluate physical activity, state of anxiety, and sleep quality are practical and easy to apply, but their effectiveness also depends on the truthfulness of the informants. Another limitation is our reliance on SDI data published by the Government of Mexico City, requiring trust in the data quality from this secondary source. Also, it is crucial to note that the cross-sectional design hinders causal inference, underscoring the need for future longitudinal investigations. Nevertheless, we were able to provide a comprehensive overview of the associations between metabolic syndrome, sleep disorders, the consumption of some nutrients, and contextual social development data such as quality and available space in the home, educational access, access to social security and/or medical services, durable goods access, sanitary adequacy, and electricity access. Moreover, as data balancing techniques continue to evolve, a variety of methods are emerging. However, in this study, we addressed only two of the most frequently used methods, ADASYN and SMOTE. It is important to highlight that we conducted only internal validation for our methods, emphasizing the necessity for external validation in larger populations in future studies.

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Informed Consent Statement: Informed consent was obtained from all subjects involved in the study.

**Data Availability Statement:** All relevant data are contained within the article. The original contributions presented in the study are included in the article/supplementary material; further inquiries can be directed to the corresponding authors.

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