Dataset Balancing in Disease Prediction

Vincenza Carchiolo^{Da} and Michele Malgeri^{Db}

Dip. Ingegneria Elettroica Elettronica Informatica (DIEEI), Università di Catania, Via Santa Sofia 64, Catania, Italy

Keywords: Machine Learning, Data Analysis, Health Informatics.

Abstract: The utilization of machine learning in the prevention of serious diseases such as cancer or heart disease is increasingly crucial. Various studies have demonstrated that enhanced forecasting performance can significantly extend patients' life expectancy. Naturally, having sufficient datasets is vital for employing techniques to classify the clinical situation of patients, facilitating predictions regarding disease onset. However, available datasets often exhibit imbalances, with more records featuring positive metrics than negative ones. Hence, data preprocessing assumes a pivotal role. In this paper, we aim to assess the impact of machine learning and SMOTE (Synthetic Minority Over-sampling Technique) methods on prediction performance using a given set of examples. Furthermore, we will illustrate how the selection of an appropriate SMOTE process significantly enhances performance, as evidenced by several metrics. Nonetheless, in certain instances, the effect of SMOTE is scarcely noticeable, contingent upon the dataset and machine learning methods employed.

1 INTRODUCTION

The importance of machine learning (ML) in healthcare is increasingly evident and significant. ML models can analyze large amounts of data, such as medical images, vital signs, and medical histories, to assist physicians in the early and accurate diagnosis of diseases. This can lead to better outcomes for patients, as it allows for the timely and precise identification of conditions. Furthermore, through the analysis of patient data, advanced customization of treatments is possible. Indeed, ML can help develop personalized treatment plans, taking into account individual variations in biological data, test results, and treatment responses, thereby improving the effectiveness of care. Finally, machine learning plays a central role in disease prevention, since ML models can identify risk factors for specific medical conditions and help prevent diseases through the early detection of predictive signs and the implementation of preventive interventions.

Medical datasets often suffer from imbalance, a critical issue for predictive modelling. When applied to imbalanced datasets, models may exhibit a bias toward predicting the majority class, resulting in Classification Bias. This bias can lead to Inaccurate Performance, particularly for underrepresented classes,

where the model fails to learn effectively from those examples or may suffer from overfitting. Balancing the dataset is thus paramount for building accurate disease prediction models. It directly impacts the model's ability to generalize correctly and make accurate predictions across all disease classes. Without proper balancing, models may struggle to generalize from the training data to new instances, impairing their predictive performance. In medical applications, dataset balancing is one of the most significant problem for several critical reasons, with the primary concern being patient safety. There are many characteristics regarding dataset balancing that become significant, such as disease prevalence, particularly when studying the class of rare diseases which might be underrepresented.

Existing literature is considered in Section 2, while the datasets are introduced in Section 3. In Section 4, the methods and results are discussed in detail. Moreover, a comparative study is presented to point out the appropriateness of the results with respect to several metrics. Finally, we consider further works and concluding remarks in Section 5.

2 DATA SET BALANCING

As commonly acknowledged, there are numerous methods for balancing a dataset. In this section, we

^a https://orcid.org/0000-0002-1671-840X

^b https://orcid.org/0000-0002-9279-3129

discuss balancing methods for classification and provide an overview of related work in the literature on methods to balance datasets, particularly focusing on health-related research. In many scenarios, the classes of interest, such as those related to rare diseases or clinically significant events, can be significantly underrepresented compared to control or normal classes. This is challenging specially during the training of machine learning models, as models tend to be influenced more by the majority class, thereby overlooking the minority class. Consequently, the model's ability to generalize to new data and correctly identify positive cases in the minority class may be compromised. Therefore, it is crucial to carefully address the issue of data imbalance in health-related datasets to ensure the construction of accurate and reliable models.

There are numerous techniques available for balancing datasets, but in this article, our focus will be on SMOTE (Synthetic Minority Over-sampling Technique) (Pradipta et al., 2021). SMOTE is one of the most widely used methods for addressing the issue of class imbalance in datasets, especially when there is a significant under-representation of minority classes compared to others. This technique is commonly applied in machine learning contexts, including classification models used to predict diseases, frauds, or other rare events. SMOTE operates based on three main components: 1. Minority Definition: this component identifies the minority class in the dataset, which is characterized by having fewer examples compared to the other classes; 2. Generation of Synthetic Examples: this step involves generating synthetic examples of the minority class. These examples are created by linearly combining nearby samples in the feature space; 3. SMOTE Procedure: for each example in the minority class, this procedure selects some of its nearest neighbors and creates new synthetic examples through a linear combination of the feature values.

Finally, by adding these synthetic examples to the dataset, SMOTE increases the amount of data available for the minority class, thus helping to balance the dataset. In addition to SMOTE, several other methods address the issue of class imbalance in datasets. Among them, we mention the following methods. The table below (Table 1) reports a comparison among them. Each technique has its advantages and disadvantages, and the choice depends on the specific characteristics of the dataset and the problem being addressed. In some cases, experimenting with different approaches may be effective in determining which one works best for the specific case.

Several authors propose various approaches to address class imbalance and feature selection prob-

lems in Clinical Decision Support Systems (CDSS). In (Sreejith et al., 2020) the authors introduce a framework that balances the dataset at the data level and employs a wrapper approach for feature selection, utilizing Chaotic Multi-Verse Optimization (CMVO) for subset selection. Performance evaluation using the arithmetic mean of Matthews correlation coefficient (MCC) and F-score (F1) indicates competitiveness of the proposed framework. Paper (Xu et al., 2021) presents a cluster-based oversampling algorithm (KNSMOTE), which combines Synthetic Minority Oversampling Technique (SMOTE) and kmeans clustering. This algorithm identifies "safe samples" from clustered classes and synthesizes new samples through linear interpolation, effectively addressing class imbalance. In a different study (Li et al., 2021; Xu et al., 2020) SMOTE is highlighted as a successful method with practical applications, alongside the introduction of a novel oversampling approach called SMOTE-NaN-DE, which improves class-imbalance data by generating synthetic samples. Additionally, a hybrid sampling algorithm named RFMSE, combining M-SMOTE and Edited Nearest Neighbor based on Random Forest, is proposed to enhance sampling effectiveness. Jakhmola and Pradhan in (Jakhmola and Pradhan, 2015) propose an interactive algorithm allowing users to customize preprocessing requirements, yielding higher quality data suitable for correlation and multiple regression analysis, as demonstrated on a diabetes dataset. Finally, in (Khushi et al., 2021) are introduced research investigates class imbalance techniques for lung cancer prediction, employing various methods including under-sampling, over-sampling, and hybrid techniques. Evaluation metrics, such as AUC, reveal the superiority of over-sampling methods, particularly random forest with random over-sampling, in predicting lung cancer presence.

3 DATASET DESCRIPTION

To analyze the balance issue in the healthcare domain, we will leverage five diverse datasets. These datasets differ significantly in terms of the number of features, observations, and imbalance ratio. Despite these differences, they all revolve around predicting medical situations through binary classification tasks. Given the inherent imbalance in these datasets, our objective is twofold: Evaluate the performance of predictions when using the imbalanced dataset and assess the impact of preprocessing techniques on preliminary dataset balancing to enhance prediction performance.

Method Advantages Disadvantages			
SMOTE Preserves information from the minority class, reducing the risk May introduce noise in the synthetic data, especially if f	he data		
of data loss. Can improve the generalization of the model. distribution is complex. Could require more computation	distribution is complex. Could require more computational time		
compared to other methods.			
Random Undersampling Simple and fast to implement. Can reduce the training time on May lead to loss of important information in the majorit	y class,		
very large datasets. increasing the risk of under-representation.			
Random Oversampling Simple to implement. Can improve the accuracy of models on May lead to overfitting if not used cautiously, especially we	vith ex-		
imbalanced datasets. cessive replication.			
Cluster Based Oversam- Effective when minority class examples form distinct clusters. Re- Requires careful parameter tuning and can be computation	ally ex-		
pling duces the risk of generating synthetic data in inconsistent regions. pensive.			
Tomek Links Enhances class separation without adding noise. May not be effective in complex class distributions.			
ENN Can improve model performance by reducing misclassification. May excessively reduce dataset size, potentially losing in	portant		
information.			
SMOTE-ENN - Combines benefits of both techniques, enhancing class separa- Computationally intensive, particularly on large datasets.			
tion and mitigating overfitting risks.			
ADASYN More effective in complex and non-uniform data distributions. Requires more computational resources compared to SMC	DTE.		
Random Oversampling with Simple to implement. Can enhance model performance on imbal- Risk of overfitting if replication is excessive, especially o	n small		
replacement anced datasets. datasets.			
Cost-Sensitive Learning Improves model performance on imbalanced datasets without syn- Requires careful weight selection and may not be universe	ally ef-		
thetic data addition. fective.			

Table 1: Comparison of some Imbalanced Data Handling Methods.

We will define the Imbalance Ratio as the proportion between the number of examples in the minority class and the number of examples in the majority class. This ratio provides a quantitative measure of the degree of class imbalance within each dataset. For example, If there are 100 negative examples (majority class) and 20 positive examples (minority class) the imbalance ratio will be 20/100 = 0.2. Obviously, the more imbalanced the dataset, the closer this value to zero.

The first dataset we discuss, Wisconsin Diagnostic Breast Cancer (WDBC) (Repository,), is the wellknown dataset that collects data for breast cancer prediction. Since breast cancer is the most common cause of cancer deaths in women and is a type of cancer that can be treated when diagnosed early, prediction is a very important aspect. This dataset has been extensively studied in the literature (Elter et al.,), which is why it is utilized in this paper. The dataset is from the University Hospital of California and can be downloaded from both the UCI Machine Learning Repository and Kaggle. It consists of 569 samples and 33 features, computed from a digitized image of a fine needle aspiration (FNA) of a breast mass and related to some characteristics of each cell nucleus (e.g., radius, texture, perimeter, area, etc.). Some of these features are more selective and decisive than others, and the determination of these features significantly increases the success of the models, which is why Feature Selection is applied to select them.

The second dataset, also widely referenced in literature, is the Heart Failure Clinical Records dataset (Chicco and Jurman, 2020a). Cardiovascular diseases (CVDs) are the leading cause of death globally, claiming approximately 17.9 million lives each year, representing 31% of all deaths worldwide. Heart failure, a common occurrence resulting from CVDs, is the focus of this dataset, which comprises 12 features aimed at predicting mortality associated with heart failure. Many CVDs are preventable through addressing behavioral risk factors such as tobacco use, poor diet, obesity, physical inactivity, and excessive alcohol consumption via population-wide interventions. Individuals with existing CVD or those at high cardiovascular risk, often due to hypertension, diabetes, hyperlipidemia, or other established diseases, require early detection and management, where machine learning models can offer significant assistance. This dataset includes medical records from 299 heart failure patients, gathered at the Faisalabad Institute of Cardiology and Allied Hospital in Faisalabad, Punjab, Pakistan, between April and December 2015. It encompasses 13 features encompassing clinical, physiological, and lifestyle-related information.

The third dataset used is Pima Indians Diabetes Database (Sigillito,). The Pima Indians Diabetes Database is a well-known dataset in the field of machine learning and healthcare research. It contains medical data from the Pima Indian population, specifically focused on women aged 21 and above from the Gila River Indian Community near Phoenix, Arizona. The dataset includes various health-related attributes such as glucose level, insulin level, BMI (Body Mass Index), age, and the presence or absence of diabetes within a five-year period following the initial examination. This dataset is widely used for developing predictive models to identify individuals at risk of developing diabetes. Due to its large sample size and comprehensive health information, the Pima Indians Diabetes Database has been instrumental in advancing research in diabetes prediction and management. Despite its significance, the dataset also poses challenges due to its inherent class imbalance and missing data, necessitating careful preprocessing and model evaluation techniques. Its availability in the public domain has facilitated numerous studies aimed at improving diabetes diagnosis and treatment strategies, contributing significantly to the broader efforts in public health and medical informatics.

The fourth dataset is a more recent data set. The

Differentiated Thyroid Cancer Recurrence dataset (Borzooei et al., 2023) is a valuable resource in the domain of thyroid cancer research. It comprises clinical data from patients diagnosed with differentiated thyroid cancer (DTC) who underwent thyroidectomy and subsequent treatment. The dataset includes various demographic and clinical variables such as age, sex, tumor size, histopathological characteristics, treatment modalities, and follow-up information. A key focus of the dataset is to predict the recurrence of thyroid cancer following initial treatment based on these factors. Researchers utilize machine learning and statistical methods to develop predictive models that can identify patients at higher risk of recurrence, thereby aiding in personalized treatment strategies and follow-up care. Due to its specialized nature and importance in thyroid cancer management, the Differentiated Thyroid Cancer Recurrence dataset has garnered attention from researchers worldwide. However, challenges such as limited sample size and data heterogeneity need to be addressed to enhance the robustness and generalizability of predictive models derived from this dataset. Overall, it serves as a valuable tool in advancing our understanding of thyroid cancer recurrence and improving patient outcomes through tailored interventions.

Finally, the last dataset used is the Sepsis Survival Minimal Clinical Records (Chicco and Jurman, 2020b). The Sepsis Survival Minimal Clinical Records dataset is an essential and widely used dataset in sepsis study and research, a severe medical condition caused by a systemic inflammatory response to an infection. This dataset contains clinically relevant and minimal information about patients with sepsis, including demographic data, vital signs, laboratory test results, and treatment information. Its simplified structure makes it particularly suitable for developing predictive models of sepsis survival and for evaluating clinical management strategies. Thanks to its availability and focused nature, the Sepsis Survival Minimal Clinical Records dataset has significantly contributed to the understanding of sepsis and the research of effective clinical interventions to improve outcomes for patients with this severe medical condition. However, it is important to consider limitations and potential biases in the data to obtain accurate and generalizable results.

The common feature shared by the aforementioned datasets is the presence of only two classes. The main data of the five datasets are summarized in Table 2, demonstrating varying numbers of features, observations, and Imbalance Ratios.

4 EXPERIMENT AND DISCUSSION

We implemented the following 10 supervised algorithms. 1. Logistic Regression (LG) is a machine learning method used for binary classification problems. Its principle of operation is based on estimating the conditional probabilities that an instance belongs to one of the two classes. It uses the logistic function (or sigmoid function) to transform a linear combination of features into a value between 0 and 1, representing the estimated probability. This value is then used as a threshold to assign the instance to one of the two classes. 2. Support Vector Machine (SVM) operates by seeking to find the optimal hyperplane of separation between classes in the case of binary classification. The separation hyperplane is defined as the hyperplane that maximizes the margin between the nearest class instances, which are called support vectors. **SVM** can effectively handle datasets with many features, and it tends to generalize well to test data, reducing the risk of overfitting. 3. Gaussian Naive Bayes (GNB) is based on Bayes' theorem and assumes that features are independent and follow a Gaussian distribution. 4. Decision Tree DT recursively splits the dataset into subsets based on the value of features, aiming to maximize the purity of each subset in terms of class labels. 5. Random Forest **RF** is an ensemble learning method that builds multiple decision trees and combines their predictions through voting or averaging. 6. Extra Tree (ET) is similar to RF but introduces additional randomness in the feature selection process. 7. K-Nearest Neighbors (KNN) operates by classifying an instance based on the majority class among its k nearest neighbors in the feature space. The distance metric (e.g., Euclidean distance) is used to measure the similarity between instances. 8. Hist Gradient Boosting (HGB is a boosting algorithm that builds a series of decision trees sequentially, each one correcting the errors of its predecessors. It uses histogram-based techniques to speed up training. 9. Bagging Classifier (BC) is an ensemble learning method that trains multiple models on bootstrap samples of the dataset and combines their predictions. It reduces variance and improves stability. 10. Finally, Multilayer Perceptron (MLP) is a type of artificial neural network consisting of multiple layers of interconnected neurons. The selection of these methods is intended to facilitate experiments showcasing the diverse impacts of various smoothing techniques. Through these experiments, we aim to ascertain the complexity of asserting a universally superior smoothing method. As we will observe, the efficacy of a particular smoothing method, which may

Dataset	# Instances	# Features	Imbalance Ratio	Feature Types	
				# Numeric	# Symbolic
Breast Cancer Wisconsin (Diagnostic)	699	9	0.59	9	0
Hearth failure	299	12	0.94	12	0
Pima Indians Diabetes Database	768	8	0.54	8	0
Differentiated Thyroid Cancer Recurrence dataset	383	16	0.39	6	10
Sepsis Survival Minimal Clinical Records	137	3	0.21	3	0

Table 2: Dataset Information.

excel in certain scenarios, could result in inferior outcomes compared to the unbalanced dataset in other cases.

Furthermore, we will employ four distinct oversampling techniques, commonly utilized in addressing imbalanced datasets, which will be referred to throughout the remainder of the paper as *Smote1*, *Smote2*, *Smote3*, and *Smote4*.

BorderlineSMOTE with 20 neighbors=20 (Smote1) is a variant of the SMOTE algorithm that generates synthetic samples only for those minority class instances that are misclassified or lie near the decision boundary (i.e., borderline instances). It generates synthetic samples by selecting a minority class instance and finding its k nearest neighbors. It then selects one of these neighbors randomly and generates a synthetic sample along the line segment joining the original instance and the selected neighbor. By setting m_neighbors=20, it specifies the number of nearest neighbors to consider when generating synthetic samples.

BorderlineSMOTE with *neighbors* = 10 and sampling_strategy =' minority' (Smote2) is a variant of BorderlineSMOTE also generates synthetic samples near the decision boundary between the minority and majority classes. Additionally, it adjusts the sampling strategy to focus on the minority class by specifying sampling_strategy='minority'. By setting $m_neighbors = 10$, it specifies a different number of nearest neighbors to consider when generating synthetic samples compared to the previous variant. SMOTE with 10 neighbors (Smote3) is a popular oversampling technique that generates synthetic samples by interpolating between existing minority class instances. It selects a minority class instance and finds its k nearest neighbors. It then selects one of these neighbors randomly and generates a synthetic sample along the line segment joining the original instance and the selected neighbor. By setting $k_neighbors = 10$, it specifies the number of nearest neighbors to consider when generating synthetic samples. SMOTE with sampling_strategy =' minority' and neighbors = 10(Smote4), similar to Smote3, generates synthetic samples by interpolating between existing minority class instances. It further adjusts the sampling strategy to focus on the minority class by

specifying *sampling_strategy* =' *minority*'. By setting $k_neighbors$ = 10, it specifies a different number of nearest neighbors to consider when generating synthetic samples compared to the previous variant. Both BorderlineSMOTE and SMOTE aim to address class imbalance by oversampling the minority class.

To assess the impact of balancing, we will utilize Accuracy and AUC. Accuracy is a general measure of the model's precision and represents the percentage of instances classified correctly out of the total instances, it's calculated as the ratio of the number of correct predictions to the total number of predictions made, and it is particularly useful when classes in the dataset are balanced but can be misleading in presence of unbalancing. AUC measures the model's discriminative ability, i.e., its ability to correctly classify positive examples as positive and negative examples as negative. All the figures are on a logarithmic scale and the blue bar refers to the analysis of the imbalanced dataset, while the others refer to datasets obtained with the four balancing methods

The result of Breast_Cancer accuracy and Auc score are shown in figure 1, let us note that while some datasets yield consistent accuracy values across methods, others exhibit significant variability depending on the method used. For the Breast_Cancer dataset, the highest accuracy (see figure 1a) values are achieved with the ET and HGB methods, whether smoothing is applied or not. The maximum score of 0.973684 is obtained for ET when no balancing is performed and for HGB when Smote1 is applied. The maximun Auc score, 0.974206, (see figure 1b) it obtained for HGB with Smote1. This, considering that AUC is less prone to overfitting, allows us to affirm that smoothing allows us to gain an advantage, albeit small

The result of *Heart_failure* accuracy and AUC score are shown in fig 2. The accuracy values reach 1 (see figure 2a) using various methods, but almost always with smoothing. Note that this data set consistently performs quite well in terms of accuracy (the worst value being 0.829268). Analyzing the results for AUC (see figure 2b) allows us to make the same considerations and therefore its study is not particularly significant for our purposes.

For the third dataset (Pima), all methods perform















Figure 2: Heart failure scores.

better with appropriate balancing, both in terms of accuracy and AUC score 3. In this case, several

methods with SMOTE allow achieving an accuracy of 0.892857 (3a). The figure 3b clearly shows that the best AUC (DT or ET method) is consistently obtained with datasets that have had SMOTE2 applied (0.856522). This result allows us to conclude that for this dataset, characterized by an imbalance ratio of 0.54, balancing is often beneficial.



Figure 3: Pima scores.

For the fourth dataset, the Differentiated Thyroid Cancer Recurrence dataset (see figure 4), the highest accuracy value (0.961039) was achieved using the DT method with Smote1 (refer to figure 4a). Interestingly, all methods improved with balancing, which is significant considering this dataset has a higher imbalance ratio than the previous three, making the impact of balancing generally beneficial. The AUC analysis further distinguishes the methods, confirming the most effective solutions (refer to figure 4b). The top AUC score is 0.945455, obtained using the DT method with Smote1.

Finally, for the fifth dataset (the most imbalanced), the behavior is nearly equivalent for any method, as adopting the appropriate smoothing method (not always the same) 5. The best choice allows achieving a value of 0.892857. A particular situation occurs for the MLP method, which performs poorly using any of the four balancing methods. In terms of AUC score, the results are still extremely diverse depending on





ex10⁻¹ 3×10⁻¹ L^R S^{IC} G^{NB} D^T R^F E^I K^{NN} H^{EB} B^C M^R

(a) Accuracy.



the method and smoothing used, much like with accuracy. However, it should be noted that for this dataset, which is the most imbalanced one, balancing can be crucial as it allows us to achieve the best result. At the same time, if used inadequately, it can even worsen the results. In terms of ACU score, the best score is obtained with DT and ET using Smote2.

The positive effect of smoothing is better appreciated by analyzing AUC, which is less influenced by overfitting compared to accuracy. For datasets where the choice of method substantially alters accuracy values, the impact of data balancing can be significant. The figures from 1b to 5b summarize the variation of accuracy and AUC values for each dataset and method. Therefore, it can be stated that there is no single most effective Smote method, but rather, the (method, Smote) pair yielding better performance should be sought.

5 CONCLUSION

A detailed analysis was conducted on five distinct datasets, utilizing various machine learning techniques to assess the impact of data preprocessing (Carchiolo et al., 2022). For each dataset, we explored the behavior of ten distinct algorithms, each with its own characteristics and tuning parameters. In order to assess the impact of data balancing, we performed the analysis both with and without data balancing techniques, considering four different smoothing approaches to handle the presence of underrepresented classes. The results obtained highlighted a significant variation in model performance based on the different combinations of machine learning method, data balancing, and smoothing techniques. It became clear that the choice of machine learning method and the application of balancing strategies must be closely integrated to achieve optimal results. In particular, we observed that while data balancing can significantly improve model performance on heavily imbalanced datasets, inadequate implementation could lead to inferior results. Furthermore, we recognized that parameter optimization for datasets characterized by imbalance requires a particularly careful and targeted approach, as the specific dataset characteristics can significantly influence the effectiveness of proposed solutions. While various approaches exist, comparing them can be challenging due to numerous tuning parameters and variations within articles. However, ongoing research suggests the importance of exploring diverse classifiers and imbalance techniques, including deep learning models, to enhance prediction capabilities and address imbalance issues effectively. In conclusion, our analysis underscored the importance of carefully considering the specific context of each dataset and adopting a flexible and targeted approach to address the issue of data imbalance in machine learning contexts. From the analysis conducted, it emerged that in the vast majority of cases, the solutions' accuracy and AUC with the application of balancing are better. Nonetheless, as future work, the analysis will be extended to a greater number of datasets and balancing methods. Another activity for future work concerns the application to datasets that involve non-binary classification to analyze whether balancing is advantageous in this case as well. Finally, precision and recall analysis could be conducted to add further confidence in the quality of the results.

ACKNOWLEDGEMENTS

The work is partially supported by UDMA project, CUP: G69J18001040007.

REFERENCES

- Borzooei, S., Briganti, G., and Golparian, M. e. a. (2023). Machine learning for risk stratification of thyroid cancer patients: a 15-year cohort study. *European Archives of Oto-Rhino-Laryngology*.
- Carchiolo, V., Grassia, M., Malgeri, M., and Mangioni, G. (2022). Co-authorship networks analysis to discover collaboration patterns among italian researchers. *Future Internet*, 14(6).
- Chicco, D. and Jurman, G. (2020a). Machine learning can predict survival of patients with heart failure from serum creatinine and ejection fraction alone. BMC Medical Informatics and Decision Making, 20(16).
- Chicco, D. and Jurman, G. (2020b). Survival prediction of patients with sepsis from age, sex, and septic episode number alone. *Sci Rep*, 10:17156.
- Elter, M., Schulz-Wendtland, R., and Wittenberg, T. The prediction of breast cancer biopsy outcomes using two cad approaches that both emphasize an intelligible decision process.
- Jakhmola, S. and Pradhan, T. (2015). A computational approach of data smoothening and prediction of diabetes dataset. In *Proceedings of the Third International Symposium on Women in Computing and Informatics*, WCI '15, page 744–748, New York, NY, USA. Association for Computing Machinery.
- Khushi, M., Shaukat, K., Alam, T. M., Hameed, I. A., Uddin, S., Luo, S., Yang, X., and Reyes, M. C. (2021). A comparative performance analysis of data resampling methods on imbalance medical data. *IEEE Access*, 9:109960–109975.
- Li, J., Zhu, Q., Wu, Q., Zhang, Z., Gong, Y., He, Z., and Zhu, F. (2021). Smote-nan-de: Addressing the noisy

and borderline examples problem in imbalanced classification by natural neighbors and differential evolution. *Knowledge-Based Systems*, 223:107056.

- Pradipta, G. A., Wardoyo, R., Musdholifah, A., Sanjaya, I. N. H., and Ismail, M. (2021). Smote for handling imbalanced data problem : A review. In 2021 Sixth International Conference on Informatics and Computing (ICIC), pages 1–8.
- Repository, U. M. L. UCI Machine Learning Repository: Mammographic Mass Data Set. http://archive.ics.uci. edu/ml/datasets/mammographic+mass.
- Sigillito, V. National Institute of Diabetes and Digestive and Kidney Diseases, note = Donor of database: Vincent Sigillito (vgs@aplcen.apl.jhu.edu) Research Center, RMI Group Leader Applied Physics Laboratory The Johns Hopkins University Johns Hopkins Road Laurel, MD 20707 (301) 953-6231 (c) Date received: 9 May 1990.
- Sreejith, S., Khanna Nehemiah, H., and Kannan, A. (2020). Clinical data classification using an enhanced smote and chaotic evolutionary feature selection. *Computers in Biology and Medicine*, 126:103991.
- Xu, Z., Shen, D., Nie, T., and Kou, Y. (2020). A hybrid sampling algorithm combining m-smote and enn based on random forest for medical imbalanced data. *Journal* of Biomedical Informatics, 107:103465.
- Xu, Z., Shen, D., Nie, T., Kou, Y., Yin, N., and Han, X. (2021). A cluster-based oversampling algorithm combining smote and k-means for imbalanced medical data. *Information Sciences*, 572:574–589.