

A Machine Learning Approach for Real Time Prediction of Last Minute Medical Appointments No-shows

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Abstract: A no-show is when a patient misses a previously scheduled appointment. No-shows cause an impact in the healthcare sector, decreasing efficiency. When a patient misses an appointment the clinic resource are wasted, postpones his or her chance to get treated for a medical condition and denies medical service to another patient. In this research, machine learning techniques are used to find patterns in healthcare data and make no-show predictions. A no-show prediction model is proposed to integrate machine learning techniques into a model that supports the testing of predictions on different datasets. The model is integrated into an online medical appointment booking platform to allow the models and predictions made, to be saved and integrated into a real-time system. Machine learning techniques are tested using three datasets with different characteristics. Through these tests, the model proposed can find the best features, which are similar in every dataset. The results obtained are compared to other prediction algorithms and techniques.

1 INTRODUCTION

A no-show is when a patient misses an appointment that was previously scheduled. This phenomenon happens in all sorts of areas, where there is the need to schedule patients or clients into a time slot. In this paper, we focus on the healthcare area. No-shows cause an impact in every hospital and clinic in the world. Attenuating the effects of no-shows in the healthcare area is something that can provide many economic and social benefits. When a no-show occurs there are two consequences, the first happens to the patient who misses the appointment who postpones his chance to be treated for a medical condition. The second one affects the hospital and other patients because there are other patients who could have used that opportunity to be seen by the doctor. This lost opportunity also means a loss of revenue to the clinic or hospital.

Given the current high demand for healthcare, wasting available resources is unacceptable, contributing to the increase in the list of patients waiting to receive assistance. To attenuate some of these consequences it is important to figure out what makes patients miss their appointments and, whether or not there are identifiable patterns that allow us to

know how likely are patients to miss their appointment.

In order to predict no show probability, the appointment data stored by hospitals and clinics around the world can be used. Using this data and combining it with machine learning techniques it is possible to find some of these patterns and obtain a probability for how likely is a patient to no-show. If these probabilities are high then specific actions can be performed by the hospital, like scheduling another patient for that time slot or contact the patient to try to confirm the appointment.

1.1 Objectives

The objective of this research is to improve and keep developing a no-show prediction system for a company that intends to incorporate it in a scheduling system for hospitals and clinics. The goal of this system is to help clinics and hospitals mitigate the negative effects of no-shows. There are three main features of this system: The first one is to notify the patients of the appointments and to confirm their presence. The second one is a prediction algorithm that uses machine learning techniques and will return the probability of no-show for all appointments. Finally, if the system detects a high probability of no-

show it will automatically try to reschedule another patient to that time slot. This research is focused on the second one by testing and improving the current machine learning techniques with real data from hospitals and clinics. To achieve this goal this paper focus on three main objectives:

- Create a prediction model that automatically returns the no-show probability for an appointment, and is able to efficiently test data from many clinics and hospitals without the need of an advanced user to tweak the system.
- To test many machine learning techniques and find out which provide the best and most consistent results across many datasets.
- To find out which features are more important to obtain better results, which prediction algorithm works best and how accurate can the predictions be in large datasets from real-world clinics and hospitals.

This work is expected to give a solid foundation to allow the continuation of tests in the no-show prediction system and give some answers to what results are possible to obtain from these machine learning methods to tackle the no-show problem.

This work is structured as follows. Section 2 includes a review on related work including previous work developed with similar goals and its limitations. Section 3 presents the prediction model developed. Section 4 describes the results that were obtained while comparing different techniques. Finally, section 5 concludes the work.

2 RELATED WORK

No-shows are estimated to have a big financial impact on hospitals and clinics and as such many studies can already be found on analysing the impact of no-shows (Neal et al., 2005), how to best deal with them and using new ways to try to predict them. All of this to reduce the level of impact they have in hospitals and clinics worldwide.

Many of these studies try to pinpoint what are the major causes of a patient no-show (George and Rubin, 2003), whether they are involuntary or not. They also focus on looking at what are the best practices to reduce the impact of no-shows, this is normally done by overbooking (LaGanga and Lawrence, 2007), but has discussed in these articles this practice may have some impact on waiting time and client satisfaction. Machine learning is a technology that has been emerging and being used in several fields and, as such, some articles studied how

to take advantage of this to reduce no-shows (Turkcan et al., 2013; Alaeddini et al., 2015; Rinder, 2012; Ferreira, 2019).

2.1 Causes of No-shows

Finding causes for no-shows is a good starting point to check if these causes can be prevented from happening and, whether or not, they can be used as ways of predicting no-shows.

Many studies give a lot of emphasis on finding out what are the causes for a patient to not show to an appointment. Missing an appointment can be a voluntary or an involuntary act, this last one being when the patient did not intend to miss the appointment. There are many reasons for not showing to an appointment these include forgetting the appointment, other competing priorities or conflicts, and the patient's health status.

The most common reason is when a patient forgets the appointment (Neal et al., 2005), for this, many clinics have already implemented a phone or e-mail reminder, which is reported to reduce no-shows (Leong et al., 2006; Liew et al., 2009). Other reported reasons for no-shows are the health of the patient which may feel better and not need the appointment anymore, other priorities like a work schedule change or having to take care of another family member and some scheduling problems due to bad quality of the service can lead to wrong appointment information and to problems in cancelling the appointment. The weather can also be a factor if it is raining or snowing people prefer to stay home and if the health problem is not serious they can no-show (Norris et al., 2014). Financial problems and lack of transportation were also some of the reported reasons.

2.2 Features for No-show Prediction

To be able to predict whether a patient is going to no-show to an appointment, it is required to have access to many factors about the appointment and the patient, which in conjunction leads to a prediction that can be stronger by having access to many factors and many similar cases. Many studies tackle this problem in an attempt to make their prediction algorithms better (Turkcan et al., 2013; Alaeddini et al., 2015; Elvira et al., 2017; Daggy et al., 2010; Huang and Hanauer, 2003). So there is already some information to help figure out which features in the appointment data of a clinic or hospital have more relevance to predict a no-show.

These features can be divided into two categories: some are relevant to the patient like gender, age,

marital status and insurance status. The others are relevant to the appointment like the day of the scheduled appointment, the amount of time between the day the appointment was scheduled and the actual appointment day and the type of clinic.

The feature found, relevant to the patient, which most articles conclude as having the most predictive power is age, where younger patients seem to no show to more appointments than other age groups (Lee et al., 2005). In the patient category, other features have some impact. These are being unmarried, not having health insurance, the severity of the illness and the scholarship level. The gender of the patient was considered by most articles as having very little impact, in other words, there are no differences between men and women regarding their attendance to previously arranged appointments (Turkcan et al., 2013). In the appointment features, it is found that the waiting time has a larger impact (Dantas et al., 2018; Norris et al., 2014). Other characteristics that also have an impact are the hour of the day, whether it is the patient first appointment (Bennet and Baxley, 2009), the medical speciality chosen, the hospital centre, whether it is a weekday or weekend, the type of appointment and the distance to the clinic. Beyond all these features there is another one that has a huge impact in predicting accurately if a patient will no-show, which is the prior no-show history, whether the patient has missed the last scheduled appointments (Dantas et al., 2018; Norris et al., 2014).

2.3 Last Minute Medical Appointments No-show Management Previous Research

There is already previous research that addressed the goal of predicting last-minute no-shows in healthcare and also contributed to the development of this prediction system. One of the first research to address this specific problem was developed by Daniel Sousa (Sousa and Vasconcelos, 2020), which focused on developing the algorithm to predict the no-shows and creating a model to replace patients that have a higher chance of not showing. There is another research by Inês Ferreira (Ferreira and Vasconcelos, 2019), which focused on testing other prediction algorithms and also created a model to send notifications to patients and update their respective no-show probability.

3 PREDICTION MODEL

This research proposes the creation of a prediction model to automate the process of pre-processing datasets and obtaining predictions from different datasets. This prediction model allows data to come from the API connected to the no-show prediction system and from CSV files. This was done to allow the system to be tested with real data from hospitals and clinics, since due to data protection measures is not possible to merge it in the API, in these early stages of development.

The prediction model has two main models and one model that supports the other two and provides a configuration file for the user. These models are discussed in the following subsections.

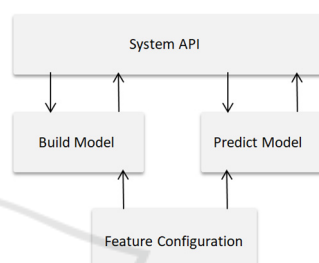


Figure 1: Models and their interaction in the Prediction model.

3.1 Feature Configuration

Feature configuration is a support file to be used in the build model and predict model phases. The goal of this configuration file is to be able to use the datasets from different clinics and hospitals that come from CSV files, without the need to change the pre-processing code every time. This configuration file is basically a simple python file with a set of variables, which need to be filled with the names of the features of the specific dataset we are using. These variables are then used by the Build Model to automatically pre-process the dataset features into the ones that will be used to train the dataset. This list of variables one that is specific to the CSV file, where the path to the CSV should be placed. The remaining features must be matched with the corresponding feature name of the dataset that will be used.

The variables chosen for the configuration file are the most common features present in most healthcare provider's datasets and to the features that possess stronger predictive power. Other features that prove to have a strong predictive power can be added in a future stage so that they can be easily pre-processed.

Once the data comes solely from the API these configuration files will not be required.

3.2 Build Model

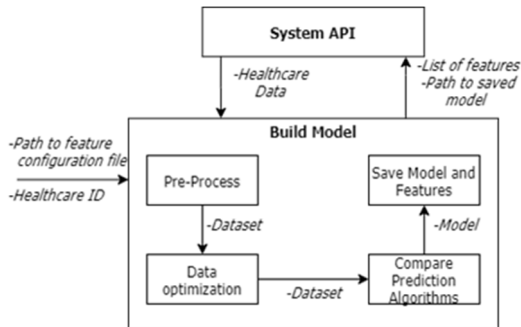


Figure 2: Build model components.

Build Model builds the prediction model, which is then used to obtain no-show predictions on the data. This data can come from a CSV file or directly from the system API. Build Model receives as arguments the path to the feature configuration file to be used and the healthcare Id of the associated healthcare provider. The model is then divided into four phases:

- **Pre-process:** Pre-processing is where the data from a specific healthcare provider is transformed into data that can be used by the prediction algorithms. This transformation adds new features from the existing ones to give the algorithms more information to learn from. It also removes or replaces missing values and transforms categorical variables with One-Hot encoding into dummy variables with values of 1 if true and 0 for false. Beyond this, all the numeric features are normalized into values in the range of 1 to 0. This allows the algorithms to learn better without giving too much weight to features with high numerical values.
- **Data Optimization:** In this phase, the data will be optimized to give the best predictions possible. The first step is choosing only the features that possess the strongest predictive power. First variance threshold is used to remove features that are almost constant since these features will not contribute to the predictions. The next step is using a feature selection algorithm. The one chosen is Boruta (Kursa et al., 2010) because it was more efficient than the other feature selection algorithms tested. To validate the features chosen by this algorithm, it is used alongside a 10 fold cross-validation and for each cross-validation fold, the features chosen are registered and only the ones that appear more than 80% of the time are chosen. The final stage is to balance the data since most of the data come imbalanced with more shows

than no-shows. This can cause the prediction algorithms to classify appointments as shows to have more accuracy. To mitigate this problem, SMOTE with Edited Nearest Neighbours (Gustavo et al., 2004) is used which balances the data by generating data samples with SMOTE and then using k Nearest Neighbours it removes those samples that are misclassified by its neighbours. After the data is balanced, it is now ready to be fed to the prediction algorithms.

- **Compare Prediction Algorithms:** It is impossible to find a prediction model that will be better for every dataset, things like the size of the dataset and the number of features can affect the quality of the predictions for some algorithms. To solve this problem, four prediction algorithms are run on the dataset on cross-validation with 3 folds only, to prevent it from being very computationally expensive. The four prediction algorithms are Artificial Neural Network, Gradient Boosting, Logistic Regression and Random Forest. These algorithms were chosen because they achieved the best results in previous no-show researches. After running the prediction algorithms, the one with the best overall score in the metric f1-score is the one chosen. This metric was chosen because it will be more important to have the right balance between recall and precision than having good accuracy. At a later stage of the no-show prediction system, the prediction algorithm that generally performs better can be chosen. This will save computation time which will be more important at that stage.
- **Save Model and Features:** After we have chosen the model it must be saved, this is done using a pickle which is a python module that allows us to save the model in a file .dat. This file can then be easily loaded to make predictions for that healthcare provider. The name of the saved file along with the features chosen in the data optimization phase will be saved in the system API to be used in the prediction phase. The reason to save the features, as well, is that the predictions need to be made with the same features the model was trained on, otherwise it will not work.

3.3 Predict Model

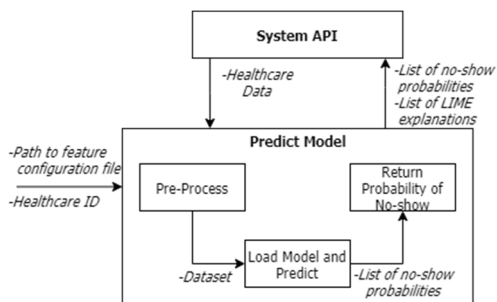


Figure 3: Predict model components.

The Predict model is the model used to obtain the probability of a patient missing his or her appointment. This model has two functions, one to obtain the probability of no-show for all the appointments scheduled, in every healthcare, and another to obtain the probability of no-show for a specific healthcare provider. The first one does not receive any argument and only works for the appointments in the API. This function is scheduled to be executed every hour or less so that the probabilities can be regularly updated and in the case, new appointments are added, we can quickly figure out what is the probability of no-show. This is especially important when appointments are scheduled for the same day or the next day.

The other function is to predict for single healthcare, this function receives as arguments the path to the feature configuration and the healthcare id. This function also works for data in CSV, and in this case, it should receive an extra argument with the appointments we want to make predictions on. The Predict model has three phases, which are:

- **Pre-process:** In the pre-processing phase, the appointments to predict are joined with the original dataset. This is done so that it is possible for the new appointments to have all the features. This is required to be able to put the right values in features like the number of appointments and the number of no-shows since this needs to be calculated for the whole data. After this, the list of features saved in the system API and associated with this healthcare provider is retrieved. The features that are not in this list of features are removed from the appointments to predict. The numeric values are also normalized using MinMaxScaler so that everything is on the same scale.
- **Load Model and Predict:** In this phase, the name of the model used to train the data is retrieved from the API and using python's

module pickle, the model is loaded. Using the loaded model the predictions for the probability of no-show are obtained for all of the appointments.

- **Return Probability of No-show:** If the appointments come from the API, the list of probabilities is uploaded to the API. Beyond this, an explanation of how the algorithm obtained that prediction is updated along with the probability of no-show. This explanation is obtained using LIME, which gives a value for how relevant the features were to the prediction and returns a list with the nine most relevant features. An example of these explanations plotted can be seen in figure 4, where the left red bars are the features that contribute to being a show and the right green bars are the features that contribute to being a no-show. With this, it is possible to have a better idea of how the prediction algorithms are obtaining those probabilities which is important in these early stages. In case the appointments come from a CSV file, the main process is the same but the results are saved to a CSV file. This file will have the appointments predicted with the original dataset features and an extra feature with the probability of no-show.

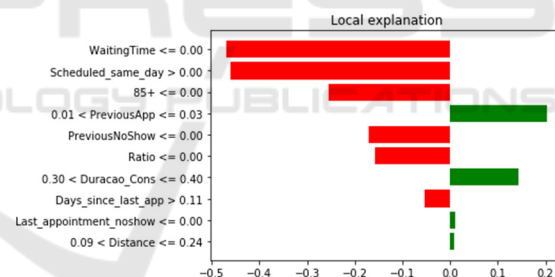


Figure 4: Plotted explanation obtained using LIME for the MD Clínica dataset.

4 RESULTS ANALYSIS

In this section, the results obtained in each of the different datasets are analysed. The prediction algorithms and machine learning techniques were tested in datasets from three different healthcare providers. With this analysis, it was possible to see which machine learning techniques and what conducts are more efficient at predicting no-shows.

4.1 Datasets Characteristics

The tests were made on three datasets, one from MD Clínica, the second one from a Brazilian dataset and the last one is from Hospital da Luz.

All these original datasets had to be transformed to be tested with the prediction algorithms. The features that were used in all datasets are the following:

- The time between the day the appointment was scheduled and the appointment itself in days.
- The number of previous appointments
- The number of previous no-shows.
- The ratio between the number of previous no-shows and the number of previous appointments.
- Whether the patient scheduled his appointment on the same day of his last appointment.
- Whether the appointment was scheduled on the same day of the appointment.
- Whether the patient's last appointment was a no-show.
- Days since the patient had his last appointment
- A binary feature for each one of the months.
- A binary feature for each one of the weekdays.
- The age divided into 10 different age groups with a binary feature for each one of them.
- Binary feature with gender.

The remaining characteristics of each of these datasets, along with the unique features in them can be seen in the following sub-sections.

4.1.1 MD Clínica

This dataset is from a dental clinic in Portugal and has 90 419 records. It comprises data from 1 January 2019 to 31 October 2019 with a no-show distribution of 65% shows and 35% no-shows.

The features unique to this dataset were distance, medic id, insurance id, part of the day and appointment duration.

4.1.2 Brazil Dataset

This dataset was obtained online from Kaggle and contains appointments from clinics in Brazil. It has 110 528 records and comprises data from 29 April 2016 to 8 June 2016 with a no-show distribution of 80% shows and 20% no-shows.

The unique features in this dataset are neighbourhood, scholarship, hypertension, diabetes, handicaps and SMS received.

4.1.3 Hospital da Luz

This dataset is from a private hospital in Portugal and has 494 627 records. It comprises data from 2 January 2017 to 31 December 2017 with a no-show distribution of 90% shows and 10% no-shows.

The unique features in this dataset are distance, medic identifier, insurance identifier, part of the day, district, speciality, whether it is the first appointment or a follow-through and whether the appointment was scheduled in person.

4.2 Prediction Algorithms

This section compares four prediction algorithms to find out, which ones can provide more reliable predictions. The four algorithms used are Artificial Neural Network (ANN), Gradient Boosting (GB), Logistic Regression (LR) and Random Forest (RF). To compare the prediction algorithms, Boruta was chosen as the feature selection algorithm. Boruta was chosen because it uses the least amount of features while achieving the same results as the other feature selection techniques tested. Since most of these datasets are imbalanced, a sampling technique for balancing the datasets was used. This will allow the prediction algorithms to find more no-shows and increase recall at the cost of some precision. The sampling technique chosen was SMOTE with Edited Nearest Neighbours because this technique achieved the best recall and f1-score than the other sampling techniques tested. This algorithm was executed in a 10 fold cross-validation and the average scores for each one of the prediction algorithms were obtained. The results obtained for each dataset can be seen in figures 5, 6 and 7, each one corresponding to a different dataset.

No prediction algorithm was found to be better in all the scenarios but the most consistent one is Gradient Boosting. The ones with more precision are Random Forest and Gradient Boosting while Logistic Regression and Artificial Neural Network have more recall. No prediction algorithm will be discarded with the tests made, as larger datasets or different features can change the type of results, this is especially the case for Artificial Neural Network which needs many data and computational power to learn efficiently.

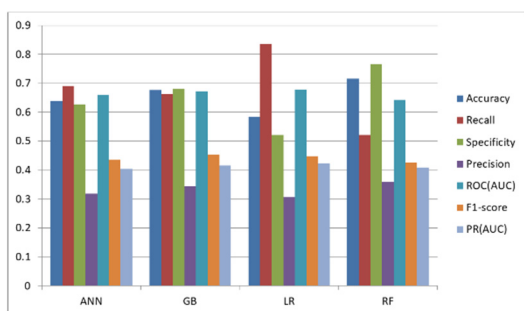


Figure 5: Results achieved by the prediction algorithms on the dataset from Brazil.

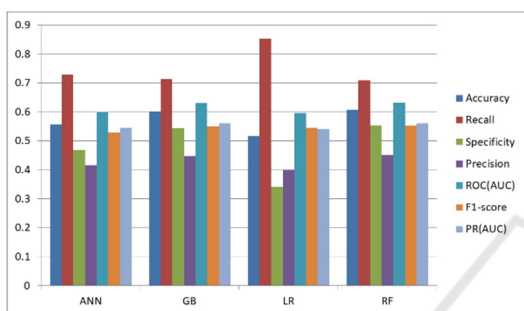


Figure 6: Results achieved by the prediction algorithms on the dataset from MD Clínica.

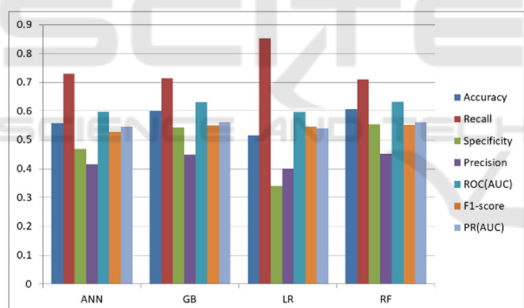


Figure 7: Results achieved by the prediction algorithms on the dataset from Hospital da Luz.

4.3 Predicting Last Week of Dataset

This comparison was done using the last week of each dataset for testing and the rest for training. A threshold of 70% was also used, what this means is that only no-shows with a probability of 70% or more are considered no-shows. This is an attempt to mimic a real-life scenario and find out how many no-shows and misclassifications happen. The sampling technique used was SMOTE with Edited Nearest Neighbours and the feature selection was Boruta. In the next table 1, we can see the comparison between the confusion matrices for all datasets and prediction algorithms. In all of the datasets, it is possible to see

that on average 50% of no-shows are found by the prediction algorithms. The prediction algorithm that finds more no-shows in all datasets is Artificial Neural Network but it also has the highest number of false positives. On the other hand, we have Random Forest with the least number of false positives but the least no-shows found, which translates to a more conservative and precise approach.

In MD Clínica dataset we can see that on average for every no-show found there is one false positive. The prediction algorithm with the best results here is Gradient Boosting since it finds almost as many no-shows as Artificial Neural Network but at a much smaller cost of false positives. The Random Forest algorithm could also be used for a more conservative approach, as it has the least amount of misclassifications, making it the most precise of the four.

In Brazil dataset, for every no-show found there is slightly more than the double of false positives. The prediction algorithms with the best results here are Logistic Regression and Gradient Boosting with similar results.

In the dataset from Hospital da Luz, there is almost the triple of false positives compared to no-shows found. It is possible to see that the more the original dataset was imbalanced the more false positives are to be expected. The best prediction algorithm here is Gradient Boosting since it is even more precise than Random Forest and finds more no-shows. Also, the number of no-shows found is not that distant from Artificial Neural Network but with less false positives.

Table 1: Comparison of confusion matrices for all datasets and prediction algorithms. The format is [TP FP][FN TN].

	MD Clínica	Brazil	Hospital da Luz
ANN	[373 385] [292 873]	[2285 5109] [1785 12808]	[222 704] [250 4335]
GB	[365 327] [300 931]	[1981 4266] [2089 13651]	[183 422] [289 4617]
LR	[309 355] [359 903]	[2088 4469] [1982 13469]	[204 697] [268 4342]
RF	[306 265] [356 993]	[1755 4051] [2315 13866]	[143 428] [329 4611]

4.4 Feature Importance

After comparing the feature importance in each of the datasets we can see that most of the chosen features are similar. This means some constant features are better at predicting no-shows. The most relevant feature is waiting time, it seems the time from when

the appointment was scheduled to the time of the appointment is crucial to find no-shows. Another very important feature is the distance, which has even slightly more importance than waiting time in the dataset from Hospital da Luz. This feature is the distance between the postal code of residence and the hospital.

Other relevant features are chosen in all the datasets, which means these features are also very important to accurately predict a no-show.

Some unique features of some datasets that got a considerable value of importance are appointment duration (Duracao_Cons), which is specific to the appointments from MD Clínica and whether a message was received (SMS_Received), which is specific to the appointments from Brazil. These features can lead to better results in the predictions and, as such, an attempt should be made to make this available on other datasets.

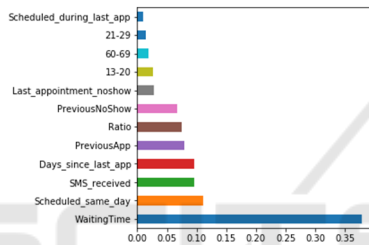


Figure 8: Feature importance graph for the dataset from Brazil.

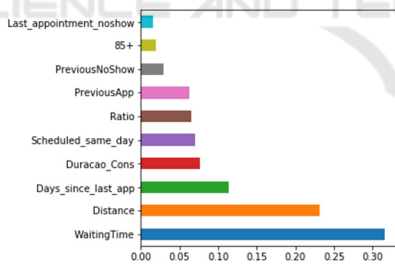


Figure 9: Feature importance graph for the dataset from MD Clínica.

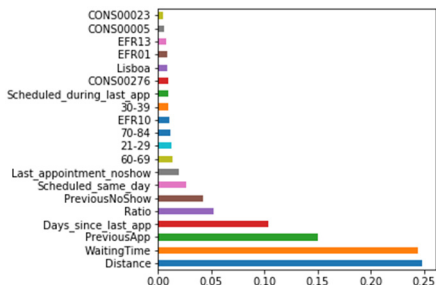


Figure 10: Feature importance graph for the dataset from Hospital da Luz.

5 CONCLUSIONS

This research was done in the healthcare area focusing on the no-show problem. It seeks to find and implement a solution capable of reducing no-shows and subsequently increase efficiency in the healthcare providers. The three major contributions of this research are next discussed.

The major contribution is the creation of a prediction model to optimize and automate testing. A prediction model was created to make the training of new models and obtaining of predictions from datasets easier and more efficient. Since all the datasets come with different characteristics and features, it would be required to change the code every time. This way the pre-processing phase and training phase were optimized, requiring a configuration file only to train the model and to make predictions. The prediction model was also integrated into an online medical appointment booking platform which is provided through an API.

Many new features were also added and tested in an attempt to figure out which features are more relevant and improve prediction results. Beyond this, machine learning techniques were tested on different datasets, in an attempt to find the techniques that perform better overall.

The main conclusions that can be made come from the results obtained. The first thing that can be concluded is that the size of the dataset did not have a large impact on results. What impacted more was the type of features available and how much imbalanced the dataset was.

The most important features are similar in every dataset and the features that were considered more important to identify no-shows are waiting time and distance. Since all these datasets were imbalanced, sampling techniques were used to counter this problem. Using a sampling technique allowed the prediction algorithms to find a much larger number of no-shows, higher recall, but at the cost of being less precise. Whether more precision is required or more recall will depend upon the clinic or hospital strategy. Some hospitals and clinics will want to keep waiting time to a minimum and favour precision, while others with less volume of patients might prefer higher recall. Having a patient confirmation strategy working alongside the no-show prediction system will be very important to reduce many of these false positives.

In the case of the prediction algorithms none of them stands out but the one with more consistent results overall was Gradient Boosting.

The results obtained are far from ideal and more features will be required to make these predictions better. We conclude that these predictions can help but are not still strong enough as a standalone strategy and should be combined with other scheduling strategies like patient confirmation.

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