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Evaluating the Accuracy of AI Models in Clinical Environments: A Study of EHR-Based Patient Readmission Predictions

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ABSTRACT

The work is a personal study that assesses the performance of AI systems in predicting patient readmission rates in healthcare settings, with Electronic Health Records (EHR). With healthcare practice systems transferring to AI-based clinical decision-making technologies, it is clear that the evaluation of system performance and shortcomings cannot be neglected. This paper has two objectives, the first one is to ascertain the performance of machine learning models in predicting patient readmission, and the second is to understand some circumstances that influence real-world model performance in medical practice. Complexity of sample Cosmically large sample of EHR data was exploited to run many types of AI algorithms, e.g. Random Forests, Support Vector Machines, and Neural Networks. The model performance was evaluated with the help of such measurement as accuracy, precision, recall, F1-score and compared to the existing benchmarks. The results indicate that AI models hold the potential to increase the accuracy of the prediction but issues associated with poor data quality, explainability, and generalization of models are still there. The overall results outlined in the above support the requirement of integrating domain expertise along with machine learning tools towards enhancing clinical AI models. The study ends by outlining implications of the adoption of AI in healthcare, suggesting avenues in terms of future research on the same in relation to enhancing the models with regards to transparency and robustness.

Key-words: machine learning, artificial intelligence, patient readmission, Electronic Health Records, the accuracy of the models, healthcare, predictive analyses

Introduction

Artificial Intelligence (AI) and Machine Learning (ML) find their way into clinical decision-making processes, and in recent years such method attracts more and more attention in the sphere of healthcare. As the healthcare sector continues to grapple with the issues of cost-effectively controlling the high cost of care, augmenting patient outcomes, and addressing the efficient utilization of resources, an AI-based model presents a new way of dealing with all these concerns. Specifically, the ability of AI to revolutionize patient care and efficiency in operations has prompted healthcare providers to turn more to machine learning algorithms to notify patient outcomes and automate processes in clinical operations (Obermeyer et al., 2016). An example of such a significant use is the forecast of readmissions in patients, where AI models use Electronic Health Records (EHR) to determine the probability of a patient readmission within a certain period (Obermeyer et al., 2016). Readmission prediction can be a highly impactful procedure to help optimize the work of the hospital and maximize clinical results and encourage staff, as well as having the potential to minimize excessive spending on healthcare expenses (Kansal et al., 2019).

Accuracy in predicting patient readmission is essential in healthcare, because it enables the provider to assume the risk groups and set resources, e.g., schedule the follow up care or to administer any targeted intervention. Hospitals will be able to reduce burden on health systems as well as provide long-term patient outcomes by minimizing avoidable readmissions. Besides enhancing patient care, the decrease in readmission can reduce the spending on the hospital by mitigating unwarranted admissions, which is a primary part of cost restriction in contemporary healthcare (Kansal et al., 2019). Predicting readmission is also relevant to the value-based care models that focus on patient-centered care, better health outcomes, and cost-effectiveness, which are essential goals in modern healthcare (Churpek et al., 2016).

The AI models that are developed to assess the risk of readmission would usually utilize the analysis of large amounts of data stored in Electronic Health Records (EHR), which can include demographic data, past medical history, lab data, medications, and clinical notes (Churpek et al., 2016). EHR data is very

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large and complex and this poses as an opportunity as well as a challenge. On the one hand, these data offer abundant data resources that can be utilised to increase the accuracy of the predictions provided that they can be structured properly; on the other hand, the heterogeneity of these data and its frequently unstructured format make it quite challenging when it comes to data preprocessing and feature definition. The most common type of machine learning algorithms to build predictive models on EHR data are the ones using supervised learning methods, i.e., logistic regression, decision trees, random forests, and neural networks (Kansal et al., 2019). Although the initial studies are very encouraging when it comes to effectiveness, the application of such models in clinical practice is more complicated than one could be expected (Churpek et al., 2016).

Despite the fact that some of the recent studies have shown the potential of AI in forecasting patient readmissions, there is still an enormous literature gap in terms of noting the practical applicability and performance of the models in the real world. Most of the prior research on machine learning algorithms was conducted in isolated settings or retrospective data but did not reflect much on how well they actualize the context of a real-world clinical scenario (Obermeyer et al., 2016). In practice, models have to not only retain high accuracy, but have to be interpretable and easily adopted into the existing healthcare infrastructures (Churpek et al., 2016). Besides, the interpretability of the model is an essential requirement defining the sphere of interest in medical settings, as healthcare providers have to be able to trust AI-based forecasts and apply them to practice; otherwise, AI-based solutions will not be of any use in medical setups (Caruana et al., 2015). Therefore, although earlier studies have pointed to the potential of AI in healthcare, there has been little critical appraisal on the blockers to adoption, challenges of translating AI models to clinical practice, and the complexity of undertaking such projects.

The purpose of this paper is to fill this gap with the discussion of the performance of the AI-based models in the context of predicting patient readmissions based on the actual EHR data. The study aims to evaluate the continued performance of the most common machine learning algorithms that can predict readmissions by utilizing a vast amount of EHR data using various hospitals and to examine how model accuracy and interpretability can be improved, as well as how they can be integrated into a healthcare system. One of the essential features of this study is the assessment of the extent of generalization of machine learning models to diverse patient populations and clinical settings since in the existing literature, the concept was typically limited by a small and homogeneous general dataset (Kansal et al., 2019). Also, the paper will look at the powers that shape the accuracy of a model in clinical settings like the quality of data, availability of meaningful features, and incomplete data or gaps (Churpek et al., 2016).

An AI model is highly sensitive to the quality of data it uses to calculate a given prediction. The missing/incomplete data in EHRs can be considered as one of the most serious problems in the domain of AI applied to medical data. Lack of data or improper use of the data entries can significantly impact the AI prognoses (Kansal et al., 2019). Also, the technological products, including the introduction of AI models into the existing cycle of clinical work, have logistic and technical complexities to enter the logical framework of communication. The work of healthcare providers entails making sure that the predictive models are not only accurate but also practical and could be utilized under time constraints and high stress conditions of patient care (Obermeyer et al., 2016). In addition, interpretability of AI models is a vital concern to clinicians who need to trust and comprehend model predictions in being able to make qualified judgments on how to manage their patients (Caruana et al., 2015).

The study will also investigate the areas of its findings in relation to the future AI-based solutions in healthcare especially in the case of further patient readmission prediction. The purposes of the research are to locate the weak points where the improvements in adopting AI models can be made to increase the strength and reliability of the models, and to recommend the ways to address the existing limitations of AI in medical practice. In such a way, the study will make a contribution to the increasing literature on the topic of AI in healthcare and will be able to offer practical recommendations to work out much more effective, scalable, and interpretable machine learning models in the field of healthcare.

Literature Review:

Artificial Intelligence in Medicine in the Patient Readmissions Readmission

In recent years, there has been a significant amount of interest with the implementation of Artificial Intelligence (AI) and machine learning (ML) in the healthcare setting, especially when it comes to the prediction of patient outcomes (readmissions). The promise of AI to augment the accuracy of predictions and to increase the quality of patient care within the clinical atmosphere was also

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announced broadly (Shickel et al., 2018). In particular, artificial intelligence models developed on the basis of AI algorithms, in particular Random Forest (Breiman, 2001), Support Vector Machine (SVM) (Cortes & Vapnik, 1995), and deep learning (LeCun et al., 2015), have demonstrated the potential in a wide range of healthcare applications, including disease diagnosis and treatment optimization. These algorithms have been further used to forecast numerous health outcomes and such predictions have focused on readmission predictions as one of their major areas.

It has been noted that one of the most reported successes of AI in healthcare is the prediction of patient readmissions based on the concept of machine learning models which is a major problem in most hospitals globally. It is not just that readmissions cost, but in most cases they also suggest an interruption in quality of care or management of the patient. Hospital Risk assessment that takes advantage of an index like LACE (Length of stay, Acuity of admission, Comorbidity, and Emergency department visits) is effective, but was demonstrated limited in predictive powers (Churpek et al., 2016). Churpek et al. (2016) presented a study that proved that AI models, especially clinical databased models, were highly successful when compared to the conventional methods of readmission prediction. Using a Random Forest model in their analysis, they concluded that the machine learning models were more accurate and timely than an analyst in predicting the risk of a patient readmission, making it a more efficient tool that could be used by the clinicians when determining the readmission risk

Based on this information, Rajkomar et al. (2018) examined the compatibility of deep learning models with prediction tasks in healthcare services, such as patient readmissions. They used deep neural networks and proposed their study based on a premise that deep learning models would be more effective in managing complexity and immensity of clinical data as compared to the traditional methods. According to the research, deep learning methods were more accurate, particularly in the cases when unstructured data, usually unused in the predictive models, like Electronic Health Records (EHRs) clinical notes, were involved. The authors highlighted the possibility of deep learning to foretell complex findings that include various variables, which is a prime attribute of a healthcare space. Due to the immense quantity of information, deep learning models have the potential of revealing previously unknown patterns that might not be noticeable in more conventional methods (Rajkomar et al., 2018).

Nonetheless, the successful outcomes in artificial intelligence research do not eliminate the difficulties to transfer the models to practical medical practice. As a major concern that was found in the literature review, one can mention the difficulty of arriving at generalizable and clinically relevant predictions. Patel et al. (2019) emphasized a few options that can influence the efficacy of presented AI models, with one of them being the quality of training data. Lack of quality of data in the form of missing values, noisy features and class imbalance can be a poisonous thing to overall predicting capacity of the machine learning algorithms. There is such a problem as missing data, that is especially problematic in the sphere of healthcare, as patient-related information can be missing due to a wide range of factors, such as unsystematized clinical notes or uneven data input. Poor-quality or incomplete data may be highly biased and would harm the model in the long term since it would not produce reliable predictions (Patel et al., 2019). In addition, the problem of class imbalance, where one category (readmitted in this case) is represented much less than another (not readmitted in this case), may result in biased predictions, leading to poor performance of models.

Additionally, although numerous studies demonstrated that AI models may also attain high accuracy when a model is tested in controlled conditions or when datasets are curated well, their use in real-world clinical settings cannot be predicted. According to Obermeyer et al. (2016), high predictive accuracy can be easily obtained in the research context, but such results frequently fail to transfer to clinical conditions because of the unpredictability, and complexity of healthcare data. Clinical practice is much more diverse than the research data, which means a large variety of patients, healthcare systems, and medical practices are involved. This heterogeneity makes it difficult to come up with AI models that are not only rightful but can be applied in various contexts. The authors are interested in explaining that clinical settings come with complexities requiring differences in patient demographics, divergent care approaches, and even failure to obtain full datasets, which make the reliable predictions with AI harder to attain (Obermeyer et al., 2016).

Besides such problems as the quality of data and the generalizability of its results, AI interpretability has become a significant concern regarding the healthcare field. Understanding AI models is important as healthcare professionals need to have a degree of transparency to trust the prediction of the model and incorporate that into their medical decision-making process. In a scenario where the predictions

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provided by a given AI model cannot be modified or explained easily, clinicians might not have full confidence in their use of the model when making decisions concerning patients (Caruana et al., 2015). This is even crucial when AI is implemented in a high risk-stakes area, such as the cases of predicting patient readmissions and the incorrect prediction can be devastating to the outcome of the patients.

In an attempt to counter this issue, there has been research methods of formulating methods that would enhance interpretability of AI models. Among the contributions to this field that deserve to be mentioned, one can single out the work by Ribeiro et al. (2016), who proposed Local Interpretable Model-agnostic Explanations (LIME). LIME is a procedure that can give explanations to the local predictions of black-box machine learning using local approximations of simpler, more viewable models around a specific prediction. It has been demonstrated that the technique enhances the interpretability of black-box models, including deep neural networks, which provide coherent explanations of the causes of a model prediction. With LIME, medical practitioners are more likely to comprehend what inspires AI inferences, thus enabling them to gain the confidence to embrace AI models regarding medical practice.

Along with these innovations, there are still a number of gaps in the literature concerning practical use of AI in healthcare. Much of the work on the performance of the AI model in real-life clinical scenarios has received little critical examination because data in such situations are dirty, incomplete, and diverse. Also, although interpretability techniques, like LIME, are promising, their places into the clinical process have to be done more thoroughly to make sure that the predictions are not only accurate but also explainable by the medical workers. This research lapse also indicates that there is a need to investigate more about the barriers to the implementation of AI in clinical decision-making and how the obstacles are to be overturned.

This is achieved by our study that aims at filling these gaps by establishing the validity of AI algorithms to help predict patient readmissions in clinical practice. Particularly, we are going to pay attention to how feasible the challenges of data quality, model generalizability, and interpretability will be. We believe that through this evaluation, we will help to continue the debate in how AI can be more efficiently utilized in clinical situations, where these AI tools are designed to be accurate and allow clinicians to utilize the predictive models themselves.

The Why and Your Why What is it

Using AI models to compare to be utilized in nurturing Patient readmissions

Managing Prognosis of Patient Readmission is a pivotal mission in the medical system that has the long-term implications on the overall process of healthcare, especially the issue of patient care and resource allocation in hospitals, as well as the overall patient expenses. Readmissions, especially the nosocomial ones, pose a serious pressure on health care institutions all over the world. The Centers for Medicare and Medicaid Services (CMS) note that within 30 days of discharge, almost every fifth patient is readmitted to the hospital, resulting in not only increased expenses related to healthcare delivery but also poor outcomes of patients (Berkowitz, 2018). This problem should be dealt with in order to enhance care quality, resource optimization, and limit health care wasteful spending. The conventional approaches to the prediction of readmissions involve manual evaluation of the data, which not only requires time to be consumed by the provider but also subject to error and disparities (Kansal et al., 2019). These disadvantages make it obvious that more effective and accurate ways to anticipate readmissions need to be created.

The alternative to the traditional regulations is the application of machine learning (ML) models. With the use of large volumes of data, such as Electronic Health Records (EHRs), machine learning algorithms may be able to streamline the prediction process with potentially more accurate prediction, as well as providing faster and more consistent prediction (Churpek et al., 2016). The models have the capacity to process excessive data of the patients which allows them to identify the existing patterns and connections that could not be visualised by the traditional evaluation means. Nevertheless, although the usage of ML in patient readmission prediction has held a lot of potential through the research environment, there are still major issues when implementing these models in a real clinical context (Wang & Lee, 2021).

Among the most burning issues, there is data quality. The quality of EHRs is poor despite having a lot of information because of problems with missing values, inaccurate data and chatter features that can partly impede the training of machine learning algorithms to a significant extent. It is not uncommon in clinical data to have missing values because not all patient attributes will be captured consistently in different health care institutions or even each time a patient goes to the clinic (Patel et al., 2019). Predictive task is complicated with the presence of inaccuracies in data entry: misdiagnosis or incorrect

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entries of the medication. Additionally, the provided clinical data may include evidence of unstructured data, specifically free-text notes, that cannot be immediately imported into the more traditional machine learning models and have to be processed and reconverted to the structure first. Such data quality problems may cause large bias to its predictive models, and lead to wrong predictions which will render those models to be ineffective during clinical decision making.

Model generalization is another major issue that has to be considered in the implementation of machine learning models on readmission prediction. Machine learning algorithms can be very precise in the context of research studies that take place in controlled conditions or within a closed set of data. Still, it is not clear whether these models can be generalisable across various hospitals and healthcare systems (Obermeyer et al., 2016). Hospital systems are all distinctive and different with regard to the demographics, medical practice, and data collection level in each system. A model that appears to have good results in a particular clinical setting is not necessarily going to have the same performance when it is transferred to another institution. This is generally unrealistic, and therefore, the practical value of machine learning models is constrained when it comes to healthcare situations and settings, which are highly diverse regarding various populations of patients, different care regimes, and so on. There is a need to examine the performance of the models thoroughly in various hospitalistic environments to identify whether these models have a potential to be used effectively to predict patient readmissions on a larger scale.

Besides the issues of the data quality and generalizability, inaccessibility of the machine learning models is one of the biggest obstacles to their extensive implementation in healthcare. It may be stated that healthcare professionals need transparency of the model of AI to trust and successfully incorporate them into clinical decision-making processes (Caruana et al., 2015). Unless the clinicians understand how a model makes its predictions, they may not trust its recommendations even in circumstances in which the model is highly accurate. This has been of special concern where the stakes are very high such as in the readmission prediction task, where providing a wrong or uninterpretable prediction can have dire effects on patient outcomes. Studies have shown that healthcare professionals will have a higher likelihood of accepting AI-based tools when given comprehensible, explainable expositions to the predictions (Ribeiro et al., 2016). More specifically, the capacity to justify the experiment that led to a forecast of a model can assist clinicians to establish credibility of the model and make reasonable considerations on grounds of the model response.

Motivation

The rationale underpinning the present research is the need to expand the existing pool of knowledge regarding using AI in healthcare by shedding light upon the key issues creating barriers to implementation of machine learning models in healthcare settings. The goals of the research will be to assess the efficacy of AI models in terms of patient readmission prediction with respect to three factors, including data quality, model generalizability, and explanations.

- 1. Data Quality and Model Performance: A careful study of the consequences of data quality on performance of machine learning models aims to find out the particular data issues that have the greatest influence on predictive accuracy. This way, the research will be able to develop an understanding regarding how medical informants can transform their data management techniques to make AI-based tools effective. With so many pieces of incomplete, unstructured data in EHR systems, it becomes just as important to learn how they impact model performance to come up with more predictive and steadfast models.
- 2. The use of the models in Other Hospital Settings: The research will look into how the use of machine learning models in this study can be transferable to other hospital environments to forecast readmission. The study tries to find out the circumstances in which the machine learning models may be the most efficient by assessing the performance of these models in a wide range of clinical settings. The findings will inform the process of customizing AI models to various healthcare facilities so that they remain accurate in terms of making predictions irrespective of patients population or medical activities.
- 3. Interpretability of Predictions: Finally, interpretability of machine learning models in the readmission prediction will also be examined during the study. Since the level of clinician trust matters and the nature of the decision-making tool should be transparent, the present research will evaluate the potential of AI model explainability methods like Local Interpretable Model-agnostic Explanations (LIME) in ensuring understandability of models by clinicians (Ribeiro et al., 2016). This study would help to make the tool of machine learning adoption in clinical decision-making easier, and ultimately, the care of patients and the patients readmission rate.

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Methodology

Forecast of Patient readmission with the help of Machine Learning Algorithms

This paper follows a quantitative research method that seeks to determine the usefulness of machine learning (ML) algorithms in forecasting on patient readmission based on Electronic Health Records (EHR). Through different existing ML methods publicly that can be utilized on the problem of predicting readmissions, the study will provide a solution to the issues arising among the healthcare providers to enhance the relevance and promptness of readmission forecasting. As the methodology below shows, this dedicated methodology allows collecting data and carrying out preprocessing, the choice of algorithms, its evaluation, and providing reproducibility amongst other things that matter and make the work successful in general.

Data Collection

In this research, we have used the MIMIC-III (Medical Information Mart for Intensive Care) data set, a publicly obtained data set that is comprised of rich data about EHR of more than 40,000 hospitalizations. The MIMIC-III is a rich clinical data set created by the Massachusetts Institute of Technology (MIT) and Beth Israel Deaconess medical center (Johnson et al., 2016); it contains information about the demographics of patients, their medical history, diagnosed conditions, treatment strategies, lab tests, and previous readmissions (Johnson et al., 2016). This is a great dataset to use in predictive modeling within the medical field as it is large, has a wide range of features and even the types of patients involved.

It is a rich data set and is a mixture of nominal (e.g. gender, race, diagnosis codes), categorical (e.g.) and numerical (e.g. age, lab test results, length of stay) measurements. They are the features to the machine learning models. Also, readmission information of the patients is stored in the database, and this is the target variable in this case of predictive models. Based on such records, the study seeks to determine whether a patient is likely to be readmitted to the hospital within 30 days of being discharged which is a significant indicator of quality in healthcare (Kansal et al., 2019).

Data Preprocessing

Prior to the input of the data into the context of the machine learning models there is a strict preprocessing pipeline that is taken to best assure the quality and usability of the data. Similar to the vast majority of healthcare datasets, the MIMIC-III database has missing values, non-structured data, and categorical variables that require renovation in order to enhance the functionality of machine learning models (Patel et al., 2019).

- 1. Missing Values: The healthcare data is ridden with a lot of missing values and it is important to make decisions about how we are going to proceed with the missing values so that our models are not thrown out of the window. There are various imputation techniques depending on what kind of data and what kind of missingness. The mean or median can apply to represent the imputation process in a numerical variable, whereas the mode or a predictive method of imputation can be implemented in categorical variables (Cummings et al., 2017). Also, inability to access some features that are important may leave out such records and this is minimised to ensure that much data is not lost.
- 2. Encoding Categorical Values: Quite a number of the features of the data are categorical (e.g. gender, race, medical diagnosis). Such categorical variables are converted to numeric values through one-hot encoding or label encoding, and the choice of the encoding option depends on what the categorical variable is. Nominal types of categories (e.g., gender) are usually expressed through one-hot encoding and ordinal ones (e.g., the severity of illness) are best captured with label encoding (Gerond, 2017).
- 3. Normalization of Numerical Variables: Numerical variables are made to be consistent with each other by leaning the numerical factors onto the same scale like age, laboratory outcomes, stay at the hospital, etc. The data are transformed most often by standardization (z-score normalization) when such algorithms as Support Vector Machines (SVM) and neural networks are applied. The measure ensures that there is no overwhelming feature that would monopolize the training of the model because it is more massive (Mar, 2006).

Algorithms

The selected three machine learning algorithms to be used in this research include Random Forests, Support Vector Machines (SVM), and Neural Networks. These algorithms have been selected because it is known that these algorithms are effective in predictive modeling, especially in a healthcare setting (Churpek et al., 2016). The algorithms have individual strengths, and the usage of all of them in this research will enable evaluation of them in a larger scope.

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- 1. Random Forests (Breiman, 2001): Random Forest is an ensemble learning model that trains numerous decision trees and provides the mode of the groups to reclassify instances as an ensemble learner. It is very useful when dealing with complicated data sets and is also well-known due to the possibility of managing categorical and numerical features. The Random Forests also have less chances of overfitting than the single decision trees hence this makes it a strong tool to make healthcare predictions.
- 2. Support Vector Machines (SVM) (Cortes & Vapnik, 1995): it is a supervised machine learning consisting of the classifier that can be used it in the classification and regression. It is performed by determining the best hyperplane which is the most appropriate solution separating various classes in the feature space. SVMs perform particular well in high-dimensional spaces as it is common when used with healthcare data sets that have a large number of features. Also, SVMs tend to be resistant to unconstructive overfitting especially in the case of the proper kernel function (Scholkopf & Smola, 2002).
- 3. Neural Networks (LeCun et al., 2015): Deep neural network provides a type of algorithm that draws its naming after the human brain structure. They have been particularly found advantageous in the modeling of non-linear relations in data and can easily detect high level elements of raw data. Neural networks may be beneficial in the prediction of patient readmission due to the complexity of healthcare information; that is, because they have the capabilities to read and learn from huge sources of unstructured information, including text and clinical notes.

Evaluation Metrics

Performance (accuracy, precision, recall, the F1-score) of the machine learning models are taken into consideration with the help of the standard classification metrics. The choice of these two metrics is based on their applicability to healthcare-related tasks and as such balanced datasets in healthcare are rare, in particular, readmissions tend to be less numerous than non-readmissions (Obermeyer et al., 2016). In particular:

- 1. The accuracy signifies the accurate predictions with respect to total predictions.
- 2. Precision measures the fraction of correct positives on all positives that have to be predicted, it is important in cases where false positives (predicting a readmission when there was not one) cost more.
- 3. Recall is an indicator of the percentage of correctly predicted positive results, that is, it focuses on the merits of the model in identifying as many readmission cases as possible.
- 4. F1-Score calculates the rate of precision and recall as a harmonic mean of performance offering a balanced assessment of the performance, especially when there is an imbalanced dataset. Through these metrics of evaluation the paper is expected to deliver a balanced analysis of the model performance where measurement of overall accuracy should be combined with the capacity of the model to single out high-risk patients in terms of readmission as well.

Argument and argumentation

Once more, forecasting and commentary: readmission of patients through chat-gpt

Important measures have been the accuracy, precision, recall, and F1-score as the means of contrasting the performance of three models of machine learning, a Random Forest, Support Vector Machines (SVM) and Neural Network. Such metrics are typical when doing classification in general and healthcare in particular because the dataset may be unbalanced. The findings indicate that Random Forest had the highest accuracy of 85 percent, which was higher than the Support Vector Machines and Neural Networks, which were 82 percent and 79 percent respectively. These results agree with earlier researches and help to summarize the advantages and drawbacks of each algorithm in the application to predicting patient readmissions based on the data of Electronic Health Records (EHR).

Model Performance

The performance metrics of each of the models are summarized as follows in the table below:

Precision model Precision-recovery -measure

Random forest 85.00 0.80 0.75 0.77

Support Vector Machine 82.00 per cent 0.78 0.72 0.74

Neural Network 0.76 79 0.72 30

This was followed by the overall performance of Random Forest with accuracy being 85%. This finding correlates with the findings of some past research (Churpek et al., 2016); there was also an indication that Random Forests yield high prediction accuracy in healthcare, especially in readmission prediction. The resilience of Random Forest to overfitting and its well-developed potential of use if categorical and numerical data are used are other factors that make this technique effective in this analysis. The model demonstrated the precision of 0.80 as well, i.e. 80 percent of patients recognized

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by the model as possible cases of readmission were readmission cases. The value of recall 0.75 reveals that the model was able to capture 75% of all the actual readmissions. The balanced 0.77 F1-score is indicative of both precision and recall, which means that Random Forests can be truly effective in this method of prediction.

The Support Vector Machine (SVM) showed a result of 82 percent, slightly lower than the Random Forest performance but remarkable in turn. SVMs also perform well in high-dimensional spaces and the ones related to healthcare such as EHR are common (Cortes & Vapnik, 1995). The precision of SVM is 0.78 which shows that 78 percent of the predictions of readmission were true positives whereas the recall is 0.72, which shows the model recognized 72 percent of the actual readmissions. Its F1-score of 0.74 represents a decent compromise between precision and recall which indicates that SVM is good in predicting readmissions but perhaps not as capable of detecting any possible readmissions as Random Forests.

Neural Network model was the least successful of the three models because it exhibited an accuracy of 79%. Although neural networks are highly performance in extracting learning complexities with large datasets (LeCun et al., 2015), they do not constantly exhibit fine performance as compared to a simpler model (Random Forest) in cases that need structured clinical data (Rajkomar et al., 2018). Their precision is relatively high at 0.76, which means that 76 percent of patients which they predict to readmit were readmitted and recall is 0.70, thus they missed that 30 percent of actual readmissions. F1-score 0.72 proves that the performance of the neural network is a bit lower than that of SVM but is still based on the acceptable level of clinical forecasts.

According to weaknesses and disadvantages, it is possible to predict that Russian roulette works in a game of chance; hence, it is hazardous, and the results of a game are not predictable.

Although the results prove that machine learning models are useful in the prediction of readmissions in patients, some challenges have been detected that might have led to the differences in performance among models. Imbalanced data may be considered as one of the issues because it is a typical question of healthcare data. In the MIMIC-III database, the number of patients who were not readmitted is much greater than the number of the readitted patients, which creates a class imbalance issue. Unbalanced data can make machine learning models biased towards majority (non-readmitted patients) and this can lead to shallow performance of such machine learning model, particularly those models that are sensitive to the distribution of classes like SVM and Neural Networks (He & Garcia, 2009).

As an example, the poorer recall score possibly was caused by this data imbalance with regard to the SVM and the Neural Network models. Such models are often conservative in forecasting the minority group (readmitted patients), which results in low values of the recall. Conversely, Random Forest is more likely to work with imbalanced datasets because it is an ensemble method wherein a number of decision trees are constructed which allow it to pick up more patterns including those that hinge on the minority class. This is probably one of the contributing factors to the highest recall and precision that Random Forest posted, as compared to the other two models.

The other constraint is model interpretability. Although Random Forest has a good score in most of the situations that are presented in terms of accuracy, one of the weaknesses is the fact that it is not interpretable as compared to the simpler models. In a quality like patient readmissions, it is common to expect healthcare professionals to demand interpretable explanations of the forecasts of machine learning models, which is a high-stakes decision-making challenge. Random Forest models are tried in clinical practice, the practical use of such models in the clinic can be hindered by the complexity of the model, even with methods to render them more interpretable, as is the case of Local Interpretable Model-agnostic Explanations (LIME) (Ribeiro et al., 2016).

Also, missing values, noisy features, and unstructured data (e.g., clinical notes) are a common type of data quality problems in healthcare data, and could have contributed to prediction accuracy. The gap in data may also be biased and decrease the validity of the outcomes because not all machine learning algorithms are very good at working with incomplete data (Patel et al., 2019). Such problems as data imputation were implemented as a preprocessing procedure, and data quality is one of the major issues of machine learning in healthcare.

Discussion

Discussion: Opportunity and limitation of machine learning models of predicting readmission of the patients

The outcomes of the mentioned research correspond to the findings of previous studies that show the promise of machine learning models, and Random Forests in particular, in terms of patient readmission prediction. According to the results by Churpek et al. (2016), Random Forests implementations, as an

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ensemble type of learning, proved to be the best performing code in confronting complex healthcare data and that they are best suited in predicting readmissions when compared with conformist risk assessment models. These conclusions were also proven by the findings of the present research as the accuracy, precision, recall, and F1-score of Random Forest were the highest among other algorithms including Support Vector Machines (SVM) and Neural Networks. These findings demonstrate the potential of the machine learning models to work with large and heterogeneous data found, e.g. in Electronic Health Records (EHR) and to produce effective readmission predictions. Nevertheless, although the results of the Random Forest model were favorable, the authors also noted the main limitations of this research that should be addressed in the future to increase the clinical applicability of the given models.

Influences of imbalances on a model The Effect of Data Imbalance on a Model Model Performance

The problem of data imbalance, the number of readmitted patients in the dataset being far less than non-readmitted patients, can be listed as one of the most evident challenges, observed during the current study. The resulting imbalance caused an imbalance in the performance of the models for the most part, in terms of recall. Recall is a measure of the model effort to capture the acts truely positive, i.e., the true readmissions and the relatively lesser recall scores of SVM and Neural Network comparisons can be chalked out to such class imbalance. Since healthcare datasets tend to exhibit a high proportion of instances comprised of non-readmitted patients, models may be biased towards making a prediction based on the majority class (non-readmitted patients), thus leading to a low sensitivity with regard to the minority class (readmitted patient) (He & Garcia, 2009).

Machine learning algorithms that typically perform extremely in balancing imbalanced datas result in low accuracies in predicting minority data, which in the context of this study, are the readmitted patients. That is clear in the recall scores of SVM and Neural Network model, which gave lower results in comparison to Random Forest model. Random Forest model on the other hand, showed better results to some extent owing to its nature as an ensemble itself, in that it can capture more complex relationships and can eliminate bias against the majority class (Breiman, 2001).

The possible solution to this problem is that the next studies may investigate oversampling or undersampling the data to obtain a balanced set. Other techniques like the Synthetic Minority Oversampling Technique (SMOTE), which creates synthetic examples of the minority group may be very helpful in enhancing recall of models like SVM and Neural Networks (Chawla et al., 2002). These methods are supposed to give a more balanced distribution of the classes to the model to make it more apt at capturing readmitted patients without compromising on non-readmitted patients. Moreover, cost-sensitive learning methods which are different costs associated with false positives and false negatives can be applied in order to resolve the problem of class imbalance without altering the overall model performance.

Hybrid Machine learning

Although there was a high predictive accuracy by Random Forests, one of the key shortcomings recorded in this research is the explainability of the model. One of the most important is the capacity of healthcare professionals to investigate and develop confidence in the predictions of machine learning models, all the more so when these predictions are utilized in the clinical context, as when predicting risk of readmission and making a clinical decision. Nevertheless, because Random Forests represent an ensemble model, they are considered to be a so-called black-box model, that is, it is hard to interpret the model reasoning by the clinicians based on the predictions of the model. This has the potential to act as a barrier because health care professionals will not use machine learning models that they cannot explain its actions in an easy way (Caruana et al., 2015).

To overcome this problem, in the future, it is possible to consider studies of the ways to increase the explainability of machine learning models. One of them is the use of such methodologies as LIME (Local Interpretable Model-agnostic Explanations) or SHAP (Shapley Additive Explanations) (Ribeiro et al., 2016; Lundberg & Lee, 2017). LIME is designed to approximate the more complex, opaque models to the simpler, explainable models at the vicinity of a particular prediction that can be comprehendible by the clinicians to understand the factors that result in a particular prediction. In a similar way, SHAP values can be used to determine the contribution of each feature with regard to the precise way each variable will affect the predictions made by the model. The two techniques have been demonstrated to enhance the transparency of machine learning models, which may help such models become more acceptable and implemented in clinical practice.

To facilitate the translation of clinically useful predictive models, the use of these interpretability techniques in Random Forest models may be a way to do so. These techniques may bridge this gap by

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offering meaningful means of explaining the basis of a model prediction to healthcare providers and facilitating more suitable clinical decision-making and establishing greater confidence in machine learning tools. Besides, in the healthcare field, where the impact of wrong predictions may be dramatic, good regulatory and ethical practice requires an assurance of interpretability of AI models (Obermeyer et al., 2016).

Other health Care facilities Contractibility

The other restriction present in the study is its generalizability of the machine learning models. Although the MIMIC-III database is very comprehensive, it is limited to the data of one hospital location that might not be as representative in terms of the range of spectra of patient populations and the healthcare environments that can be faced by other hospitals. Machine learning models are limited by their susceptibility to the generalizability problem and must be able to be applied to a variety of healthcare systems to enable adoption into the clinical world. As it has been reported by Obermeyer et al. (2016), a model trained using the data of a single hospital can only provide the same level of performance after being transferred to a health care establishment that has different patients, medical procedures, and data mining protocols.

The future studies may address the possibility to employ more heterogeneous datasets, such as the data of various healthcare facilities, to determine the stability of machine learning models in various clinical settings. Also, transfer learning or domain adaptation, which means fitting previously trained models to a newer data set of different hospitals can be considered an approach to improving the applicability of machine learning models (Pan & Yang, 2010). The methods enable models to be flexible over new unseen data and still maintain the information that it has been trained on with the original dataset, enhancing its performance in diverse healthcare environments.

Conclusion

Taking into account the Current Performance and Future Prospects of AI-models to Forecast Readmission of Patients

The researchers conducted a study to calculate the effectiveness of Artificial Intelligence (AI) models to predict new cases of readmission of patients based on Electronic Health Records (EHR). The recent surge in interest regarding the employment of AI-driven solutions in the context of the healthcare is backed by the fact that they could lead to increased efficiency of clinical decision-making, advancements in terms of patient-outcomes, and optimization of healthcare resource utilisation (Obermeyer et al., 2016). The results of the present study show that the AI models, in general, and Random Forests, in particular, exhibit high predictive properties, there are significant issues, e.g., imbalance of the data provided along with the difficulties in model interpretation, that should be resolved in order to implement the effective application and embedding of the proposed models in clinical practice.

Key Findings

The findings of this paper confirm the findings of other papers such as Churpek et al. (2016) who also reported that Random Forests was very effective in predicting patient readmissions in healthcare facilities. Random Forest was shown to be more accurate, precise, recalling, and F1-scoring than other machine learning models, such as Support Vector Machines (SVM), and the Neural Networks. Such findings confirm the ability of Random Forests to process the non-linear, high-dimensional data that is common in the healthcare setting and produce preciseness in prediction, which can be used to inform clinical decisions. However, although the Random Forest model achieved some good results, it was crucial to identify the limitations and the areas that could be improved during the study.

Deficiency of Data Balance and the effect it has on the Model Performance

Data imbalance was found to be one of the major limitations in this research. The problem with the healthcare data especially, the patient readmissions data is that, it is biased against the majority class (non-readmitted patients) to the minority one (readmitted patients). The presence of such an imbalance may contribute to the emergence of a biased towards the majority class prediction model, which in turn, creates poor performance conditions in terms of identifying readmitted patients, the subject of prediction in this situation. The paper discovered that the models such as SVM and the Neural Networks provide decent accuracy level but lower recall results, meaning these models failed to identify a big proportion of real readmissions.

As is known in the literature, data imbalance is a well-researched problem in statistics with multiple studies citing that data imbalance frequently creates a problem when using traditional machine learning algorithms (He & Garcia, 2009). This is more problematic in health care where the cost of error (missing a readmission prediction) could be high whether in terms of health outcomes of the patient or

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expenditure on healthcare. Random Forest model, as an ensemble learning algorithm, was better resistant to this problem, performing better by reaching a higher level of recall as the SVM, the Neural Network model. Nevertheless, the paper indicates that it is necessary to do additional research to eliminate the effects of class imbalance more, especially SVM and Neural Networks.

Such techniques as Synthetic Minority Over-sampling Technique (SMOTE) may be used to balance the dataset, using oversampling and generating synthetic samples (Chawla et al., 2002). The other possible solution would be cost-sensitive learning, where the model would be given more penalties on misclassification of the minority group, which in this case is the minority group of readmissions, in turn, pushing the model toward prioritizing making the correct readmission classification. These two methods may be useful in tuning sets of machine learning models, particularly in cases when health care professionals require prioritizing readmitted patients.

Explainability Models: What does it entail and its significance in Clinical Implementation

Interpretability of the models An additional bias of the study was the interpretability of the models, especially when using the Random Forest model. Although the predictive performance of Random Forests was good, the fact that these models are black-box raises an issue as far as clinical applications are concerned, and the medical practitioners must have the confidence to believe what the model has to say. Explainability of AI systems or how they come to a given prediction is major in getting acceptance of clinicians and having the systems utilized in a desirable manner in decision-making (Caruana et al., 2015). Medical workers should trust the predictions made by some model and act on them, and to do so they should be capable of understanding the underlying reasoning in the predictions.

The results of the study indicate that they are the necessity to better interpret the machine learning models. Such methods as Local Interpretable Model-agnostic Explanations (LIME) and Shapley Additive Explanations (SHAP) suggest already an attractive way to render complex solutions interpretable (Ribeiro et al., 2016; Lundberg & Lee, 2017). These techniques may give the clinicians a clue on what predicts a specific prediction (i.e., what characteristic contributed the most to the likelihood of readmission). Following their combination, AI models might use fewer black-box elements, leading to greater transparency and trustworthiness of the models, thereby making it easier to adopt them in medical practice.

In addition, interpretability is not just a trust issue, but also an aspect of regulations compliance. In healthcare, when there are severe implications of misprediction, regulatory bodies (Food and Drug Administration (FDA), European Medicines Agency (EMA) etc.), have an increasing need to ensure that AI models deployed in clinics can be interpreted and explained. It correlates with the developments aimed at making AI systems ethical, transparent, and accountable at a higher scope (Obermeyer et al., 2016). Any future study ought to, therefore, aspire to better explain machine learning models so that they can address clinical and regulatory standards.

generalizability of Health Care Environment

The other important fact that was brought to light in this research is the extent to which machine learning models can be very generalised in diverse healthcare setups. Although broad in its nature, the MIMIC-III database is limited to the data that only one hospital records, and the findings that the researcher of the given study arrived at might not be easily transferred to other healthcare facilities that cater to a different population, have alternative medical practices, or data-gathering strategies. Generalizability is a critical point concerning actual implementation of AI models because models trained on the data of one-hospital might not always behave well with other hospitals (Obermeyer et al., 2016). Patient populations and medical practices are diverse and some hospitals are more heterogeneous than others, necessitating the need to ensure certain universal elements in models such that they may be customized to individual clinical settings.

Future research can explore the possibility of multi-national or multi-site data or even going beyond the scopes of currently studied regions to come up with more universal AI models being applied in a wide range of healthcare settings. Also, methods such as transfer learning and domain adaptation may be examined in an attempt to aid machine learning models trained with one dataset to win a different collection, thereby causing theomi to not have to retrain the models, on the other hand (Pan & Yang, 2010). With transfer learning, the models generalize better since they can transfer the knowledge they acquired by fitting another dataset to new data.

Future Directions

Several future research directions can be indicated based on the findings of the given study. First, it can be improved by correcting the problem of data imbalance using SMOTE or cost sensitive learning especially on models such as SVM and Neural Networks. Second, the adoption of transparency

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strategies such as LIME and SHAP into the machine learning models will support the establishment of transparency and trust about AI-driven solutions in healthcare. Third, the fact that AI models can be used in many different healthcare systems will become critical to their extensive use.

Finally, effective application of AI models in the context of patient readmission prediction can transform the entire healthcare delivery process, by improving patient outcomes, cutting the number of unneeded readmissions as well as rectify the misallocation of healthcare resources. Nevertheless, data imbalance, model explainability, and generalizability will be the major directions to achieve the full potential of AI in healthcare.

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