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A Machine Learning Pipeline and Application for Automatic Classification of Clinical Documents

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Abstract---Healthcare industry has many associated services including research on various trends or patterns in diseases and patients' life style. With the emergence of Artificial Intelligence (AI), it is made possible that problems in healthcare domain can be solved by using Machine Learning (ML) techniques. One such problem considered in this paper is known as clinical document classification. Existing methods in this area lack a systematic approach in filtering out false positives. In this paper we proposed a ML framework that considers pipelining of ML models at multiple levels. In the first level, clinical documents that do not have any content related to smoking are discarded. In the second level, the documents that talk about known smoking cases are retained. In the third level clinical document are classified into two categories such as currently smoking and past smokers. We proposed an algorithm known as Learning based Clinical Document Classification (LbCDC). This algorithm makes use of three models in pipeline in order to perform classification of clinical documents at multiple levels of granularity. Our experimental results revealed that the proposed system is efficient in clinical document classification.

Keywords - Clinical Document Classification, Machine Learning, Learning Based Framework, Exploratory Data Analysis

I. INTRODUCTION

Clinical document classification is one of the research areas on healthcare domain. It has potential to improve operational efficiency and research pertaining to useful insights for decision making. Clinical document classification is useful at the level of healthcare units, state and central governments. It is also useful at global scale to understand disease and patients' life style patterns and help in making policy decisions. As the AI became popular ML models are widely used to solve real world problems. Clinical document classification is the problem that could be solved using ML models. However, existing methods in this area lack a systematic approach in filtering out false positives. There are many contributions found in the literature that are based on ML models.

Emilio et al. [3] used radiological reports for their empirical study. They used ML models to perform classification of such records useful to hospitals. Bevan et al. [6] proposed a hybrid ML model to detect cancer and come up with cancer statistics from death certificates. They used rule based phenomena for automatic detection of cancer and render its statistics. Zaid et al. [8] proposed a pipeline of deep learning models to detect cervical cancer from documents and classify the disease. Lin et al. [11] proposed a methodology based on deep learning for automatic analysis of heterogeneous medical data. Alzubaidi et

al. [14] proposed an application based on deep learning tools to study medical images. Their study could help in discovering new knowledge from the medical images. Gupta et al. [18] used deep learning to make a histopathological model for image classification for distributed environments. From the literature, it is observed that pruning documents prior to classification is important. This insight is incorporated in the proposed system of this paper. Our contributions are as follows.

- 1. We proposed a ML framework that considers pipelining of ML models at multiple levels.
- 2. We proposed an algorithm known as Learning based Clinical Document Classification (LbCDC).
- 3. We built an application for evaluation of our system. Our experimental results revealed that the proposed system is efficient in clinical document classification.

The remainder of the paper is structured as follows. Section 2 reviews prior works on clinical document classification. Section 3 presents the proposed system. Section 4 presents results of our experiments. Section 5 concludes our work and throw light on future scope

II. RELATED WORK

This section reviews prior works pertaining to clinical document classification based on machine learning techniques.

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Heath et al. [1] focused on automatic classification of health records in medical domain. Jahanzaib et al. [2] used ML models to classify electronic medical records in healthcare domain. It has modus operandi to fulfill this use case towards automation. Emilio et al. [3] used radiological reports for their empirical study. They used ML models to perform classification of such records useful to hospitals. Jonathan et al. [4] investigated on possibilities in healthcare domain with the emergence of AI based approaches. They found the utility of ML models for solving problems in medical domain. Eli et al. [5] proposed a deep learning model known as NiftyNet that is designed to analyze medical imagery. Bevan et al. [6] proposed a hybrid ML model to detect cancer and come up with cancer statistics from death certificates. They used rule based phenomena for automatic detection of cancer and render its statistics. Victor et al. [7] proposed a deep learning model that works in two stages for finding entities in medical documents and then find relationships among important medical texts. Zaid et al. [8] proposed a pipeline of deep learning models to detect cervical cancer from documents and classify the disease. Yunlu et al. [9] used deep learning based method for automatic classification of documents based on respiratory patterns in the patients' related documents. Jihad et al. [10] use clinical notes from medical domain and investigated on finding mental status of patients.

Lin et al. [11] proposed a methodology based on deep learning for automatic analysis of heterogeneous medical data. Jain et al. [12] explored different ML tools available for automatically finding useful information and solve problems in different domains. Pérez-García et al. [13] proposed a library for medical image analysis using Python language. It is developed for various ML approaches including pre-processing, sampling and classification. Alzubaidi et al. [14] proposed an application based on deep learning tools to study medical images. Their study could help in discovering new knowledge from the medical images. Tan et al. [15] proposed deep learning methodology for automatic monitoring of cardiovascular diseases in real time. Min et al. [16] used deep learning

approach for automatic detection of presence of diseases in human samples. Pandey *et al.* [17] investigated on medical imaging and analysis using ML models. They could throw light on possible challenges and opportunities. Gupta *et al.* [18] used deep learning to make a histopathological model for image classification for distributed environments. Yasar *et al.* [19] investigated on deep learning models to understand medical documents and detect cases of Covid-19. Nathan *et al.* [20] studied hematological malignancies in medical field using ML models. Different methodologies are found in [21]- [30] for medical document analysis. From the literature, it is observed that pruning documents prior to classification is important. This insight is incorporated in the proposed system of this paper.

III. PROPOSED SYSTEM

This section presents the proposed system that includes a ML framework, different ML models, dataset and performance evaluation procedure.

A. The Framework

We proposed a framework, shown in Figure 1, for automatic classification of clinical documents. The framework is based on machine learning. The framework is designed to have multiple ML pipelines. Model 1 pipeline is designed to know whether given clinical document corpus has any information related to smoking habit of patients. This pipeline is meant for discriminating the clinical documents that have some text related to patient smoking. Thus in this model, the documents that have smoking information are retained and the rest are discarded. The retained documents are used in model 2 pipeline which is designed to classify clinical documents into smoked and never smoked categories. The smoked category of documents is retained while the other documents discarded. The smoked category of documents is considered in the model 3 pipeline where the system classifies clinical documents into two categories such as presently smoking and past smoker

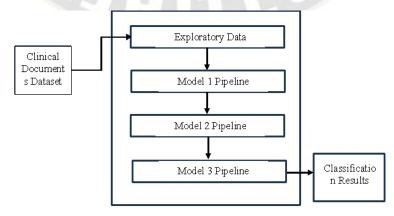


Figure 1: Overview of proposed ML based framework for clinical document classification

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After completion of all pipelines, the final document classification results contain two categories such as presently smoking and past smokers. Different ML models are used in the proposed framework. In other words, each pipeline has set of ML models such Logistic Regression (LR), Naïve Bayes (NB), Random Forest (RF) and Gradient Boosting (GB). LR is a ML model that is based on supervised learning. Eq. 1 reflects logistic model obtained from logistic function.

$$Z = \alpha + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_k X_k \tag{1}$$

The linier sum of Eq. 1 is then substitute with a function and the resultant expression is provided in Eq. 2.

$$f(z) = \frac{1}{1 + e^{-z}} \tag{2}$$

The expression is formulated as a mathematical model. The probability is modelled reflecting conditional probability as expressed in Eq. (3).

$$P(D = 1|X_1, X_2, ..., X_k) = P(X)$$
 (3)

Finally, the logistic model is constructed and expressed as in Eq. 4. The logistic model is used for classification of medical documents in the proposed framework.

$$P(X) = \frac{1}{1 + e^{-(\alpha + \sum \beta_i X_i)}}$$
 (4)

NB model is also widely used model for classification in real world applications. It is based on Bayes Theorem. In fact, it is a collection of algorithms meant for solving classification problems. The NB model is based on the expression in Eq. 5.

$$P(A/B) = \frac{P(B/A)*P(A)}{P(B)}$$
 (5)

In terms of input and output variables X and y, it is possible to rewrite the above expression as in Eq. 6. $P(y/X) = \frac{P(X/y)*P(y)}{P(X)}$

$$P(y/X) = \frac{P(X/y) * P(y)}{P(X)} \tag{6}$$

Because of the naive assumption that variables are independent given the class, we can rewrite P(X/y) as follows:

$$P(X/y) = P(x_1, y) * P(x_2, y) * \dots * P(x_n, y)$$
 (7)

The goal of NB is to detect classes with highest probability. This model is widely used in classification tasks as it has potential to discriminate class labels. Thus the NB model finally focuses on maximum y value as expressed in Eq. 8.

$$y = argmax_y[P(y) * \prod_{i=1}^n P(x_i, y)]$$
 (8)

Random Forest (RF) is yet another model that is used extensively in many ML applications. It has many underlying decision trees that form an ensemble of models to improve classification performance. Gini index is often used in RF for branching nodes in different trees of RF as expressed in Eq. 9.

$$Gini = 1 - \sum_{i=1}^{C} (p_i)^2$$
 (9)

Probability and class are used to determine to know Gini of each branch. It is also possible to exploit entropy to determine branching as expressed in Eq. 10.

$$Entropy = \sum_{i=1}^{C} -p_i * \log_2(p_i)$$
 (10)

Random forest makes use of ensemble approach with voting in order to arrive at final classification results. Gradient boosting model is also based on decision tree. The tree can be modelled as in Eq. 11.

$$h_m(x) = \sum_{j=1}^{T} b_{jm} l_{Rjm}(x)$$
 (11)

Finally, Eq. 12 and Eq. 13 provide the modus operandi of the GB model used in the proposed framework.

$$f_{m}(x) = f_{m-1}(x) + \sum_{j=1}^{T} \alpha_{jm} l_{Rjm}(x)$$
(12)
$$\alpha_{jm} = argmin_{\alpha} \sum_{x_{i} \in R_{jm}} L(y_{i}, f_{m-1}(x_{i}) + \alpha)$$
(13)

The GB model needs computation of pseudo residual, alpha and also regularization mechanism in order to get rid of overfitting.

Notation	Meaning
α and β_i	Parameters to be estimated
P(X)	Denotes $P(D = 1 X_1, X_2,, X_k)$
X	Set of variables
P(X)	Denotes probability dynamics
Argmax	Denotes argument with maximum
value	
y_i	Testing data point
fi	Return value in decision tree
p_i	Relative frequency value
Entropy	Measure used for branching of tree
Y	Actual value
Gamma	Value predicted
L	Denotes loss function
f(x)	Denotes a function

Table 1: Shows notations used in this paper

The ML models in the proposed system are based on supervised learning and their approach in the classification of clinical documents is as in Figure 2.

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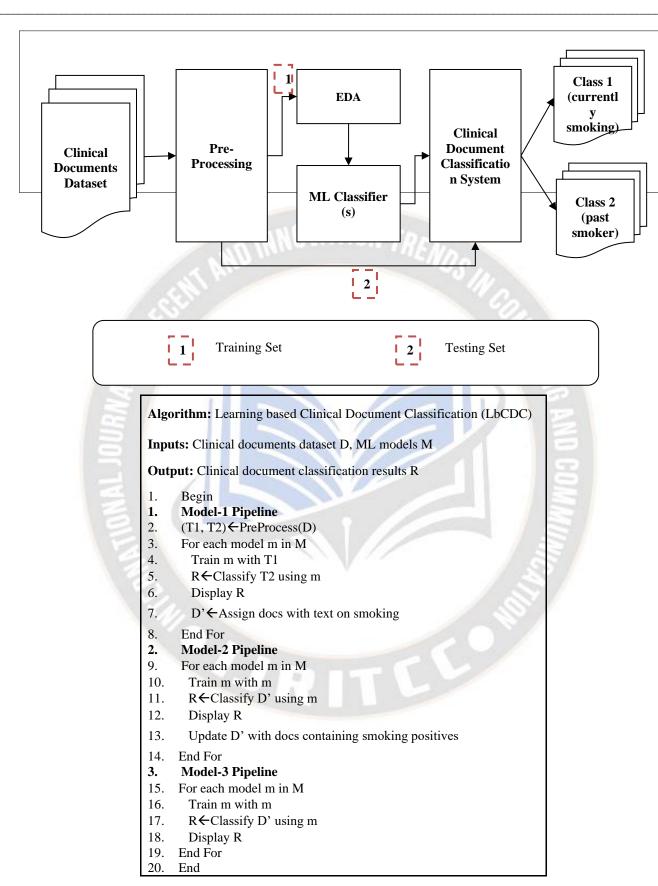


FIGURE 2: CLINICAL DOCUMENT CLASSIFICATION METHODOLOGY

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As presented in Figure 2, each classification model used in the proposed framework (for all model pipelines), the modus operandi are provided. The given dataset is subjected to preprocessing and then divided into 80% training and 20% testing. Afterwards, the training data is subjected to EDA to determine any requirement for changes in the dataset such as treating missing values. Then the training data is used by ML model to gain knowledge from the training samples. Based on the knowledge gained, the model is further used to classify clinical documents.

B. Dataset Details

Clinical documents dataset [31], known as n2c2, is collected from https://portal.dbmi.hms.harvard.edu/. The dataset contains 502 patients' discharge summaries covering various details. The documents do have smoking status which is the main focus point in this research for document classification.

C. Algorithm Design

We proposed an algorithm known as Learning based Clinical Document Classification (LbCDC). This algorithm makes use of three models in pipeline in order to perform classification of clinical documents at multiple levels of granularity.

Algorithm 1: Learning based Clinical Document Classification (LbCDC)

As presented in Algorithm 1, it takes clinical documents dataset and ML models as input and performs classification in multi-level approach. There is pipelining of models in order to filter out documents in model 1 and model 2 and finally arrive at only documents that do have details of patients with smoking habit. Then in the third model pipeline, the documents are classified into two categories such as currently smoking and past smoker.

D. Performance Evaluation

The proposed algorithm is evaluated using different performance measures obtained through the cases provided in confusion matrix as shown in Figure 3.

In this paper, the clinical document classification performance of ML models is made by comparing prediction results with ground truth. True positive indicates a positive sample is correctly detected by the proposed algorithm. True negative indicates a negative sample is correctly detected by the algorithm. False positive indicates that a negative sample is detected as positive sample by the algorithm. In the same fashion, false negative indicates that a positive sample is detected as negative sample by the algorithm.

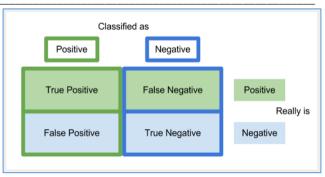


Figure 3: Confusion matrix

Precision (p) =
$$\frac{TP}{TP+FP}$$
 (14)

Recall (r) =
$$\frac{TP}{TP+FN}$$
 (15)

F1-Score =
$$2 * \frac{(p*r)}{(p+r)}$$
 (16)

$$Accuracy = \frac{TP + TN}{TP + TN + FP + FN} \tag{17}$$

Based on confusion matrix cases, different metrics are derived and used for performance evaluation. Precision, recall and accuracy are used to measure capability of ML models. These metrics result in a value between 0 and 1 reflecting least performance to highest performance.

IV. EXPERIMENTAL RESULTS

Experimental results are provided in this section for all the three model pipelines. For each underlying ML model, confusion matrix is provided as proof of performance followed by ROC curve and performance comparison of all models.

A. Model-1 Pipeline

This subsection presents experimental results in terms of confusion matrix, ROC curve and clinical document classification performance measures for model-1 pipeline.

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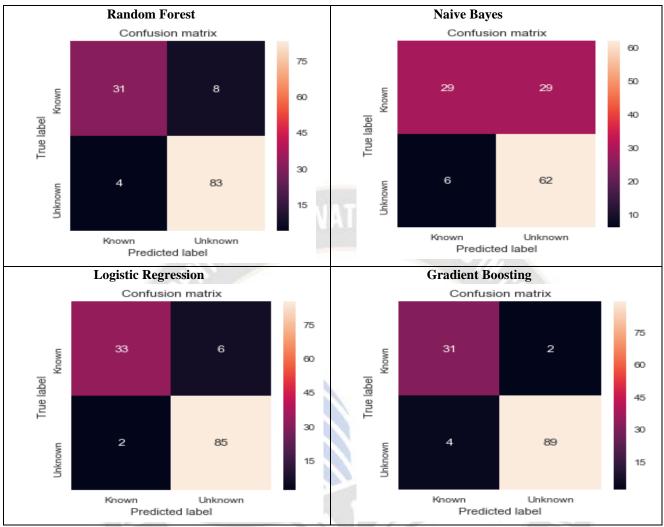


Figure 4: Model-1 pipeline confusion matrix

As presented in Figure 4, each model's confusion matrix is provided reflecting predicted and true label dynamics.

Receiver Operating Characteristic

NB, auc=0.82
— RF, auc=0.95
— GB, auc=0.98
LR, auc=0.98
False Positive Rate

Figure 5: ROC curve for model-1 pipeline

As presented in Figure 5, ROC curve provides performance of different ML models in clinical document classification.

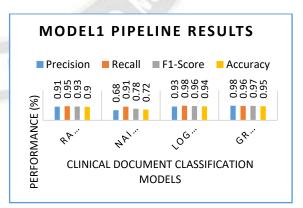


Figure 6: Shows performance of models in model-1 pipeline

As presented in Figure 6, each model is evaluated for the purpose of model-1 pipeline classification. Different performance metrics are used to evaluate the models. Higher in value for any measure indicates better performance. With

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respect to accuracy, RF showed 90% accuracy, NB 72%, LR 94% and GB 95%. From the results, it is understood that GB showed highest performance in model-1 pipeline.

B. Model-2 Pipeline

This subsection presents experimental results in terms of confusion matrix, ROC curve and clinical document classification performance measures for model-2 pipeline.

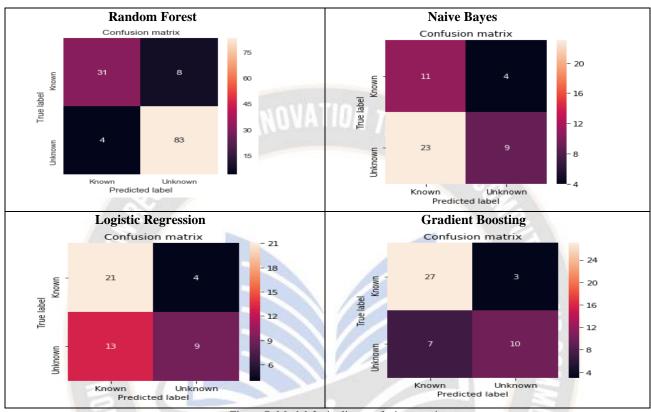


Figure 7: Model-2 pipeline confusion matrix

As presented in Figure 7, each model's confusion matrix is provided reflecting predicted and true label dynamics.

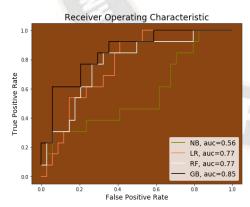


Figure 8: ROC curve for model-2 pipeline

As presented in Figure 8, ROC curve provides performance of different ML models in clinical document classification in model-2 pipeline.

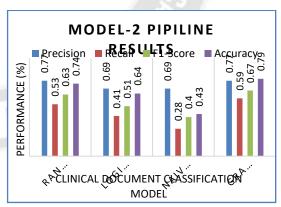


Figure 9: Shows performance of models in model-2 pipeline

As presented in Figure 9, each model is evaluated for the purpose of model-2 pipeline classification. Different performance metrics are used to evaluate the models. Higher in value for any measure indicates better performance. With respect to accuracy, RF showed 74% accuracy, NB 43%, LR

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64% and GB 79%. From the results, it is understood that GB showed highest performance in model-2 pipeline.

C. Model-3 Pipeline

This subsection presents experimental results in terms of confusion matrix, ROC curve and clinical document classification performance measures for model-3 pipeline.

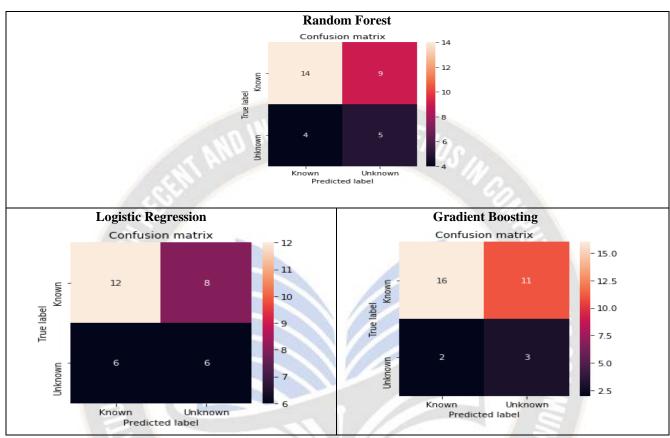


Figure 10: Model-3 pipeline confusion matrix

As presented in Figure 10, each model's confusion matrix is provided reflecting predicted and true label dynamics.

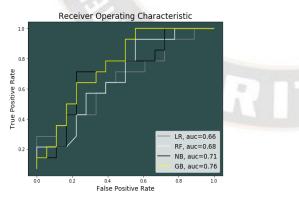


Figure 11: ROC curve for model-3 pipeline

As presented in Figure 11, ROC curve provides performance of different ML models in clinical document classification in model-3 pipeline.

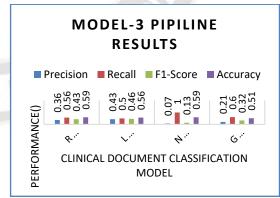


Figure 12: Shows performance of models in model-3 pipeline

As presented in Figure 12, each model is evaluated for the purpose of model-3 pipeline classification. Different performance metrics are used to evaluate the models. Higher in value for any measure indicates better performance. With respect to accuracy, RF showed 59% accuracy, NB 59%, LR

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56% and GB 51%. From the results, it is understood that RF and NB showed highest performance in model-3 pipeline.

V. CONCLUSION AND FUTURE WORK

In this paper we proposed a ML framework that considers pipelining of ML models at multiple levels. The framework is designed to have multiple ML pipelines. Model 1 pipeline is designed to know whether given clinical document corpus has any information related to smoking habit of patients. This pipeline is meant for discriminating the clinical documents that have some text related to patient smoking. Thus in this model, the documents that have smoking information are retained and the rest are discarded. The retained documents are used in model 2 pipeline which is designed to classify clinical documents into smoked and never smoked categories. The smoked category of documents is retained while the other documents discarded. The smoked category of documents is considered in the model 3 pipeline where the system classifies clinical documents into two categories such as presently smoking and past smoker. We proposed an algorithm known as Learning based Clinical Document Classification (LbCDC). Our experimental results revealed that the proposed system is efficient in clinical document classification. In future, we intend to improve our framework with deep learning techniques.

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