

UTILIZING MACHINE LEARNING TO DETECT ANEMIA

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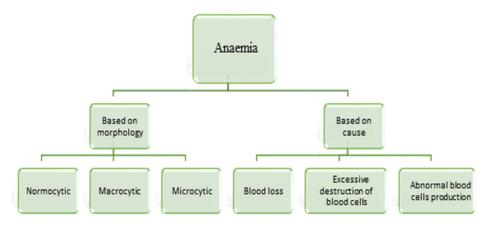
ABSTRACT

Blood is a substance that moves waste materials from the cell and carries nutrition and oxygen to it. Blood is made up of platelets, which serve to halt hemorrhaging, red blood cells, which transport oxygen, and white blood cells, which act as the body's first line of defense against infectious diseases. The most prevalent blood condition, anemia, is brought on by a deficiency in red blood cells, which prevents the body from receiving enough oxygen. Chronic anemia results from a gradual decrease in red blood cells and is frequently associated with inflammatory diseases. Acute anemia is caused by a sudden drop in RBCs. In order to identify and categories anemia, this study applies machine learning techniques such as K-nearest neighbors (KNN), Naive Bayes, and decision trees. States/UTs, Area, and other monitoring data were used as input for evaluation. Infants aged 6 to 59 months (about 5 years) with anemia (11.0 g/dl), Non-pregnant women aged 15 to 49 with anemia (12.0 g/dl), Pregnant women aged 15 to 49 with anemia (11.0 g/dl), All women aged 15 to 49 with anemia (11.0 g/dl) All ladies between the ages of 15 and 19 who are anemic : 22 (%) Males aged 15 to 49 who are anemic (13.0 g/dl) make up 22 (%), as do men aged 15 to 19 who are anemic (13.0 g/dl). The study's findings point to a wide range of potential uses for this technology in the area of medical diagnosis.

Keywords: Machine Learning, Anemia, Extraction, Identification

I. INTRODUCTION

The most prevalent blood disorder worldwide is anemia. The World Health Organization (WHO) defines anemia as a situation where there are not enough red blood cells and, as a result, not enough of them can transport enough oxygen to satisfy the body's physiological requirements. Another definition of anemia is a reduction in blood hemoglobin and hematocrit levels as well as cell bulk. Age and gender-specific numbers for normal hemoglobin and hematocrit exist. Anemia is prevalent if the values of the hemoglobin and hematocrit are below the cutoff of what is considered typical for the age and sex. Using information and resources from the 2010 WHO research on the global burden of illness, Kiassebaum et al. looked at 189 nations, both sexes, and 20 distinct age groups. They determined that 32.9% of people worldwide have anemia. Children under five and women are the most likely to experience anemia. Iron-deficiency anemia is the form of anemia that is most seen. The treatment of anemia, which has a major negative impact on quality of life and is both an illness and a



symptom of many severe diseases, can often be life-saving, making a proper evaluation the first step in treatment.

Figure. 1: Classification of anemia

Predicting anemia illness is crucial for finding other related diseases. Anemia is categorized according to its morphology or fundamental reason (Fig 1). Anemia is classified into three groups based on morphology: normocytic, microcytic, and macrocytic. Anemia is divided into three categories based on its underlying causes: blood loss, insufficient generation of healthy blood, and excessive blood cell apoptosis.

Using data gathered from the NFHS, we try to compare the effectiveness of the Naive Bayes, Random Forest, and Decision Tree algorithms in this article for the forecast of anemia illness. The reality that the disease's fundamental cause differs from region to region necessitates this study. Although the random forest classifier has previously been studied for the forecast of heart and chronic renal disease, as far as we are aware, it has not been studied for the prediction and categorization of anemia disease. This makes the product more unique.

II. LITERATURE SURVEY

Numerous data mining, deep learning, artificial intelligence, and machine learning methods have been applied to the treatment of anemia over the past ten years. The following are most prominently mentioned:

To evaluate images, [1] implements the k closest Neighbors (k-NN), support vector machine (SVM), and extreme learning machine (elm) classifiers. This method displays comparisons between the classifiers with accurate findings. Compared to k closest neighbor and support vector machines, radical learning machines are generally better.

[2] uses a machine learning method to identify anemia. The purpose of the article is to offer a remote, non-invasive, standard method that allows a fast screening to find hemoglobin levels using smartphones and AI techniques. The conjunctiva of the eye is mechanically removed as a Region of Interest from the picture of the eye after it has been photographed (ROI). Following processing, features are taken from the ROI to programme a machine-learning

system to determine whether the patient is anemic or not. The model's accuracy, precision, and memory were all 85%, and it covered 200 topics.

Based on 539 data points with 10 attributes that were gathered from labs, Mohammed Sami MOHAMMED [3], Bayesian Network (BN), Naive Bayes (NB), Logistic Regression (LR), and Multilayer Perceptron (MLP) have been used to forecast anemia. When compared to other techniques, the LR has produced superior outcomes.

Anemia has been predicted using Pooja Tukaram Dalvi, Nagaraj Vernekar [4], Decision Tree, Artificial Neural Network, Naive Bayes, and K-Nearest Neighbor algorithm. The Artificial Neural Network performs the best and K-Nearest Neighbor performs the worst among the separate classifiers. However, when used on a stacking ensemble, the classifier combo of Decision Tree and K-Nearest Neighbor gets a much better accuracy than the Artificial Neural Network. This shows that a group of classifiers performs much more accurately than a single classifier. Therefore, an array of classifiers should be used to attain the highest level of precision in medical decision-making.

In [5], This study describes the use of K-nearest neighbors (KNN), a machine learning technique, to identify and categories anemia. The study's findings point to a wide range of potential uses for this technology in medical diagnosis.

Manish Jaiswal, Anima Srivastava and Tanveer J. Siddiqui [6], In this research, we use CBC (complete blood count) data obtained from pathology facilities to explore supervised machine learning algorithms—Naive Bayes, Random Forest, and Decision Tree Algorithm—for anemia prediction. The outcomes demonstrate that the Naive Bayes method beats C4.5 and random forest in terms of accuracy.

In [7], For the forecast of anemia, supervised machine learning methods include the Naive Bayes, LR, LASSO, and ES algorithms. And assume that the patient will either be cured or uncured after 90 days (about 3 months). The outcomes indicate that the Naive Bayes method excels in terms of accuracy when compared to LR, LASSO, and ES.

Anemia and starvation are identified using machine learning, according to Richard Joseph, Vedant Sawant, and Shivani Shenai [8]. To find the different factors affecting malnutrition and connect its relationship with anemia, which has not yet been tied to existing literature, this study uses the machine learning method. Additionally, the extent to which these characteristics impact malnutrition is determined, and geographically specific characteristics are targeted for children residing in different Indian states.

Bheem Sen, Adarsh Ganesh, Anupama Bhan [9], used machine learning method to identify anemia. The automated identification of sickle cells in microscopic pictures is done in this study using image processing and machine learning methods, which then categories the RBC into three different forms: circular, elongated (sickle cell), and other shapes.

In [10], In this research, we use CBC (complete blood count) data obtained from pathology facilities to explore supervised machine learning algorithms—Naive Bayes, Random Forest, and Decision Tree Algorithm—for anemia prediction. The outcomes demonstrate that the Naive Bayes method beats C4.5 and random forest in terms of accuracy.

In [11], create a curate anaemia prediction system, three based rule categorization methods are used: ZeroR, OneR, and PART to select pertinent anemia datasets linked with "If" and "Then" procedures. In terms of the methods used, PART offered 85% more precision than

ZeroR and OneR. These methods offered a standard for other methods that were used to explain the necessary understanding of anemia data principles.

[12], utilizes the WEKA data-mining tool along with a few categorization methods, including Naive Bayes, Multilayer Perception, J48, and SMO. The J48 decision tree method provides the greatest possible classification of anemia types, according to numerous tests. This study uses a predictive model and data mining classification methods to identify the specific form of anemia in anemic individuals.

<u>N. Sasikala</u>, Gulmohamed Rasitha Banu [13], Using scientific research is conducted data extraction methods like J48 and Random Forest trees also a hoeffding tree. Consequently, the show is three categorization methods were examined for them comparison of precision using an uncertainty matrix. Random Forest tree provides greater accuracy, it has been determined. compared to the J48 and Hoeffding tree method.

Three data mining classifiers—Decision Tree, Rule Induction, and Naive Bayes—are used in [14]. The results demonstrate that Rule induction classifier performs better in predicting Hematology (Adult, Children) with accuracy of (57%-67%) than Nave Bayes classifier (accuracy of 56% for tumor of blood disease), while Decision Tree has the lowest accuracy rate for detecting the three types of diseases in our dataset.

Laengsri V, Shoombuatong W [15], A classifier model was built using five machine learning techniques: k-nearest neighbor (k-NN), decision tree, random forest (RF), artificial neural network (ANN), and support vector machine (SVM). The effectiveness of the system was evaluated and compared to thirteen pre-existing diagnostic indices and algorithms.

In this article, we analyze the prediction and categorization of anemia in patients using data mining methods, as described in [16]. Based on CBC reports and anemia severity, a decision tree for anemia categorization is created. This decision tree provides the finest anemia classification available. We have found that the C4.5 algorithm performs best and is most accurate.

<u>K. Dinakaran</u> and <u>R. Preethi</u> [17], The current research suggests a novel approach for locating possible data on blood-related diseases. Maximizing Expectations Together, the clustering algorithm and the k-Means clustering algorithm successfully group the patients based on their characteristics. It has been demonstrated that improved data yields the best results and may help the medical community identify a group of patients.

Srdjan Denic,1 Mukesh M. Agarwal,2 [18], The goal of this research was to investigate some possible factors that could have an impact on HbA2 and what that might mean for BTT screening. It is reasonable to regularly measure serum ferritin in females because iron deficiency is mostly restricted to females. Additionally, all individuals without BTT who intend to marry BTT carriers should get a Genetic test for the -thalassemia mutation. This is because marriage between two BTT carriers can be devastating for the offspring. This method of BTT screening needs to have its benefits supported by bigger research.

Thomas H. A. Ederveen, Fajar Agung Nugroho, [19], This research used a computational model to identify a causative relationship between gut microbiome profiles, RTI and systemic inflammation factors, and the efficacy of iron supplementation. The relationship between clinical factors and iron-micronutrient supplementation in infants with iron deficiency anemia was discovered through computer modelling. The biological basis of RTI, the change of the

gastrointestinal flora, and iron supplementation may all be better understood because of more targeted research.

In [20] By building a device using a Raspberry Pi that can measure the different parameters of the RBC such as area, perimeter, diameter, shape geometric factor (SGF), and detecting the central pallor and target flag, a proposed system is intended to assist medical technicians, hematologists, and pathologists in identifying RBC. This method only assists medical professionals in the early diagnosis of abnormal red blood cells; further laboratory tests must be performed to definitively identify an illness linked to abnormal RBCs.

[21], data mining methods are used to operate the system, with pattern recognition as the system's operating system. Artificial Neural Networks were able to achieve the best efficiency data mining techniques (78.31). In this manner, biochemical parameters have been demonstrated to be useful for identifying iron deficiency anemia and will assist the doctor in starting an efficient course of therapy for the patient.

<u>Kanak Meena¹</u>, <u>Devendra K Tayal²</u> [22], With the goal of lowering the risk of blood-related illness anemia, the two-methods decision tree and association rule mining have been applied and contrasted to determine which is more suitable for this specific task.

[23] In this research, there were 104 54 men and 50 women are gathered with their health data. blood hemoglobin content, anemia, and palpebral examination conjunctiva picture. For detecting anemia, the 81 samples are educated using a variety of classifiers, including Cosine K NN, Linear SVM, and Coarse Tree have been obtained. best accuracy of 82.61% in trials for Decision Tree (Coarse) 23 data.

[24], The method suggested in the current study will make it possible to identify anemia in routine clinical practice settings. A model created for this system using four distinct machine learning techniques. As categorization systems, Artificial Neural Networks, Support Vector Machines, Naive Bayes, and Ensemble Decision Tree techniques are employed. Bagged Decision Trees had the best accuracy (85.6%), followed by Boosted Trees (83.0%) and Artificial Neural Networks (79.6%).

III. METHODOLOGY

Dataset Collection:

In-depth patient data from the research was utilized in the study. As a result, we sought the advice of a qualified medical expert and identified the characteristics she used to identify different forms of anemia. The dataset's Gender, Hemoglobin, MCH, MCHC, MCV, TYPE, and Result characteristics are all utilized.

Pre-Processing:

The proposed method uses CBC test values. First, the data are pre-processed to extract. Then, we apply the Random Forest, Decision tree, KNN, Linear SVM, Logistic Regression, Radial SVM, NB classifier on it. The performance evaluation is done in terms of accuracy and mean absolute error (MAE). The mean absolute error (MAE) measures how close the predictions are to the eventual outcomes. A summary of the three classifiers findings. Accuracy has been

attained using tenfold cross-validation. The comparison of each classification algorithm's accuracy and MAE performance.

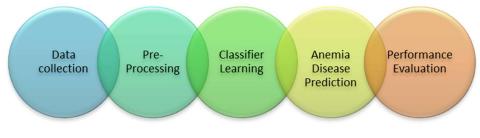


Figure. 2: Proposed diagram

Classification:

In the current study, the effectiveness of well-known classification techniques was assessed by producing an entirely new dataset. KNN, support vector machines, decision trees, and naive bayes are selected as the classifiers since they are frequently utilized in literature. These cutting-edge categorization techniques produce encouraging results. Additionally, it has been established that these techniques work well when applied to medical data. Consequently, various techniques are used, and the outcomes are contrasted.

Random forest:

The decision tree classification is the ancestor of the random forest (RF) method. It is a combination of tree predictors that employs majority polling to determine and combines the outcomes of all the trees in the collection.

Decision Tree:

A decision tree is a tree in which each leaf node represents a judgement, and each branch node

represents an option among several possibilities. It has been widely applied in many different areas. Ross Quinlan created the decision tree C4.5 (J48 in WEKA). Naive Bayes Algorithm

Naïve Bayes:

The Bayes rule of conditional probability is the foundation of the Naive Bayes algorithm. It makes use of every characteristic in the data and examines each one separately because they are all equally significant and independent of one another. Very little training material is needed.

LINEAR SVM

The term "linearly separable data" refers to data that can be divided into two groups using only a single straight line. Linear SVM is used to classify such data, and the classifier utilized is known as the Linear SVM classifier.

LOGISTICS REGRESSION

A primary use for the supervised machine learning technique known as logistic regression is classification tasks, where the objective is to determine whether a given instance will belong to a certain class or not. It is a type of statistical technique that examines the correlation between several independent factors and the dependent binary variables. It is an effective instrument for making decisions.

RADIAL SVM

When the data cannot be separated linearly, a radial kernel SVM is a useful solution. Making nonlinear modifications to the characteristics Xi to place them in a higher dimensional space is the rationale underlying creating nonlinear decision boundaries.

IV. MODEL EVALUATION METRICS

Numerous indicators are available to assess machine learning models in diverse applications. Let's examine the evaluation metrics to evaluate a machine learning model's performance. This is essential in any data science project because it seeks to estimate a model's generalization accuracy on future data.

A. Precision

Accuracy can become an unreliable criterion for gauging our success when there is a class imbalance. As a result, we also need to consider class-specific performance indicators. One of these measurements, called precision, is defined as positive predicted values.

$$PRICISION = \frac{TRUE \ POSITIVE}{(TRUE \ POSITIVE \ + \ FALSE \ POSITIVE)}$$

B. Recall

A recall is another crucial indicator; it measures the percentage of real positive cases that were accurately identified.

$$RECALL = \frac{TRUE \ POSITIVE}{(TRUE \ POSITIVE \ + \ FALSE \ NEGATIVE)}$$

C. F1-score

Precision and Recall are two significant error metrics that make up the F1 score together. In light of binary data categorization, it can be seen as the Harmonic Mean of Precision and Recall error metrics for an unbalanced dataset.

$$F1 SCORE = \frac{2 * PRECISION * RECALL}{PRECISION + RECALL}$$

D. Accuracy

An indicator of the model's performance across all classes is accuracy. When all types are equally important, it is helpful. It is calculated by \div the total number of % by the number of accurate forecasts.

$$ACCURACY = \frac{TP + TN}{(TP + TN + FP + FN)}$$

E. Mean Average Precision

It is a well-liked evaluation metric for object detection in computer vision. An instance's localization pinpoints its precise location, whereas its classification identifies its nature.

$$mAP = 1\sum_{i=1}^{n} AP_i$$

V.OBSERVATIONS, RESULTS AND DISCUSSION

We have analyzed the results from Accuracy, Precision, and Recall by comparing them with other ML algorithms.

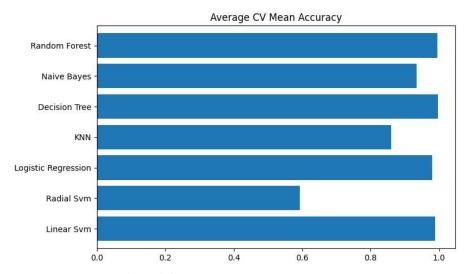


Figure. 3: Proposed Model Accuracy

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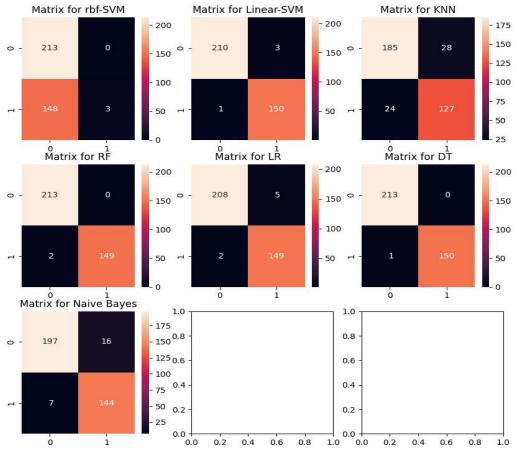


Figure. 4: confusion Matrix of classifier

| Algorithm | Accuracy | Precision | Recall |
|---------------------|----------|-----------|--------|
| Logistic Regression | 97% | 97% | 97% |
| Radial SVM | 90% | 91% | 91% |
| KNN | 80% | 81% | 81% |
| Decision Tree | 99% | 99% | 99% |
| Random forest | 99% | 99% | 99% |
| Linear SVM | 96% | 97% | 96% |
| Naive Bayes | 89% | 87% | 89% |

VI.CONCLUSION

In this study, we evaluated the accuracy of several classifiers in predicting anaemia sickness. The experimental outcome on a test dataset indicates that, when compared to Random Forest, Naive Bayes, Linear SVM, Logistic Regression and Radial SVM, the Decision Tree classification method performs best in terms of accuracy. Automatic prediction can lessen the amount of manual work required for diagnosis. In the future, automated technologies may be created to assist in suggesting additional diagnoses based on the predictions. These automated technologies may be useful in the early identification of more serious diseases. Such a disease prediction system can also suggest a treatment strategy.

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