

# Symptom based COVID-19 test recommendation system using machine learning technique

Lakshmana Rao Kalabarige<sup>a,\*</sup> and Himabindu Maringanti<sup>b</sup>

<sup>a</sup>*Department of Computer Science, GMR Institute of Technology, India*

<sup>b</sup>*Department of Computer Applications, Maharaja Sriram Chandra Bhanja Deo University, India*

**Abstract.** At present, the mankind of the entire world is under serious threat due to the unexpected COVID-19 pandemic. The advent of this pandemic exposes many drawbacks in the medical and healthcare system. As per the guidelines of WHO, the spread of the virus must be controlled through proper measures that help cease the virus. Tracing infected subjects (people/patients) is exceedingly difficult across the globe. The testing process in many countries is hampered by the unavailability of COVID-19 Test kits. Therefore, a testing process needs a robust mechanism to identify the infected subject to reduce the infection rate. To address this issue, a Symptom-based COVID-19 Test Recommendation System using Machine Learning methods is proposed and tested on real data set. It is found that the results of the system are promising and accurate up to 99%. The proposed piece of work undergoes four steps. First, it creates synthesized data set by using inputs of the Superintendent of Physical Health Centre (Rajam). Second, the synthesized data set is balanced by using Random under-sampling (RUS) followed by Synthetic minority oversampling (SMOTE). Third, different machine learning techniques such as K-Nearest Neighbor (KNN), Decision Tree (DT), Naïve Bayes, Random Forest (RF), Stochastic Gradient Descent (SGD), and Support vector machine (SVM) are applied on both the Synthesized and balanced data sets to classify subjects into different classes based on age, comorbidity-chronic disease-and other symptoms (cold, cough, fever, and breathlessness). Finally, the COVID-19 Test Recommended System is created and integrated with the best classification model. From the experimental results, it is observed that the training and testing accuracy of all the classification models is more than 99% consequently, the COVID-19 Testing recommended system also gives 100% accuracy in predicting the category of a subject based on input symptoms.

Keywords: COVID-19, machine learning, random forest, data balancing, comorbidity

## 1. Introduction

The International Committee on Taxonomy of Viruses (ICTV) labeled Severe Acute Response Syndrome coronavirus (SARS-CoV2) induced coronavirus disease (COVID-19) [1] pandemic initially started in Wuhan, China since January 2020 and spread all over the world. The COVID-19 is belonging to Coronaviridae family, and size is 65–125 nm diameter creates severe respiratory problems and ease of spread through air and droplets of human while coughing.

Currently, many developed countries in the world are severely affected by this virus due to the lack of authorized information about this virus. At the same time, ignorance, unpreparedness, and reluctance of public authorities of different countries increased the severity of this virus. The dynamic gene mutative nature of the COVID-19 virus enables it to adapt to quick changes in the environment and helps to sustain in different weather conditions. This peculiar behavior of the virus poses many challenges to the entire world. The advent of this pandemic exposes many drawbacks in the medical and healthcare system. It exposes many vulnerabilities especially related to the readiness of societies to such kinds of health emergencies. The World Health Organization (WHO) releases several guidelines

---

\*Corresponding author: Lakshmana Rao Kalabarige, Department of Computer Science, GMR Institute of Technology, India. E-mail: lakshmanarao.k@gmrit.edu.in.

to face the current situation from time to time. Many countries following either the 3T (Test, Treat, Track) or 5T (Test, Trace, Treat, Teamwork, Track, and monitoring) formula to contain the virus. Many countries have announced Lock-down to reduce the spread of the virus.

In India till 22 March 2020 around 360 people have been infected with COVID-19 and 7 were dead. At present, according to the statics of the Indian government as of 19 Jan 2021 exactly 1,05,81,837 has been infected with COVID-19, the death toll was 1,52,556, the death rate is 1.44% and discharged cases (cured) after treatment is 96.66%. This analysis shows that India is on the safer side because of strict implementation of lock-down till 31 May 2020 and subsequent unlock rules till date. On the other hand, In Andhrapradesh (AP) state 886245 people have been infected with COVID-19, 877443 were cured, active cases are 1660, the death toll was 7142, the death rate is 0.8%, and discharge rate is 99% as of 19 Jan 2021.

Mucahid Barstugan et al. [2] proposed the machine learning-based classification of COVID-19 CT images which applies feature extraction methods such as Grey Level Co-occurrence Matrix (GLCM), Local Directional Pattern (LDP), Grey Level Run Length Matrix (GLRLM), Grey-Level Size Zone Matrix (GLSZM), and Discrete Wavelet Transform (DWT) to improve the performance in classification. The Support Vector Machines (SVM) algorithm is used for the effective classification of infected regions of CT abdominal images.

Zifeng Yang et al. [3] proposed the Modified SEIR and AI prediction of the trend of the epidemic of COVID-19 in China under public health interventions which integrates machine learning techniques with Susceptible-Exposed-Infectious-Removed (SEIR) model to predict the progression of COVID-19 epidemic in china. The domestic migration and the most recent COVID-19 epidemiological data are applied to the modified SEIR model to predict the status of coronavirus in china. In Japan, according to COVID-19 policy, people with high fever for more than four days are recommended for COVID-19 test, and the people with mild COVID-19 symptoms are restricted to homes. This situation hides the deaths due to COVID-19.

The society coexisting with COVID-19 [4] discusses how badly Japan country affected due to the delay in COVID-19 testing. Hence, a recommender system for selecting a potential subject or candidate for COVID 19 testing is proposed. Machine Learning algorithms are used to classify a subject into any of the five classes such as No-symptom, General symptoms, Low-risk,

Medium-risk, High-risk. The people labeled as Low-risk, Medium-risk, and High-risk are recommended for the COVID-19 test. This proposed work considers five-cold, cough, fever, breath, and comorbidities-ill-health symptoms and age as the other important factor.

The rest of this paper is organized as follows, Section 2 discusses past literature related to different disease recommender systems. Section 3 explains the working approach of the proposed work. Performance evaluation of classification techniques discussed in section 4. Implementation of the symptom-based recommended system presented in section 5. Finally, conclusion and discussion for future scope in Section 6.

## 2. Related work

Patil et al. [5] discussed suitable machine learning algorithms used to predict, decision making, and analyze medical data related to diabetes. The authors experimented on PIMA Indian diabetic data set. The proposed method applies different pre-processing techniques, future (Forward and Backward) selection methods, and then applied five different classifiers (ADABOOST, Decision Tree, XGBOOST, Voting Classifier, and Stacking Classifier) to produce five different trained models on the PIMA Indian data set. Finally, it is concluded that the results of ADABOOST and Random Forest are best out of all five classifiers in terms of accuracy and other metrics.

Mumtaz Ali et al. [6] proposed a neutrosophic recommended system for medical diagnosis based on algebraic neutrosophic measures which identify or recommend disease of a patient based on three issues such as symptoms of a patient, the probable symptoms of a respective disease, and identification of disease based on symptoms possessed by a patient.

Chen et al. [7] proposed Disease Diagnosis and Treatment Recommendation System is an ontology-based Diabetes Medication Recommendation system which creates knowledge on different attributes – the nature of the diabetic drug, dispensing type of drug, and side effects – of each drug/medicine and ontology information on symptoms of a patient to advise/recommend a potential prescription to the diabetic diseased person. Phanich et al. [8] proposed the Food Recommendation System (FRS) which suggests a healthy and nutritional diet plan for diabetic patients. The food clustering analysis is performed by FRS to recommend food with less sugar and fat content. In recent times, the importance and usage of natural herbs for chronic diseases are increased

Table 1  
Rajam rural and urban data set from primary health center

sno	Secretariat	Number of houses	General public symptomatic	High risk factor	High risk factor	High risk factor
				General symptomatic and Age > 60	Age > 60 and comorbidities	Age <= 60 and comorbidities
1	SYAMPURAM	738	1	0	0	0
2	ANTHAKAPALLI	1271	1	0	0	1
3	G.C.PALLI	616	2	1	0	0
4	GURAVAM	983	17	9	4	1
5	SOPERU	1278	14	10	1	0
6	BODDAM	1646	12	6	11	11
7	POGIRI2	923	2	1	1	1
8	RAJAYYAPETA	1216	0	0	0	0
9	VOMMI	648	7	4	0	0
10	DRNVALSA	781	21	5	0	3
11	PENUBAKA	1161	3	1	0	1
12	POGIRI1	887	2	1	0	0
13	M.J.VALASA	1001	7	3	17	15
14	AGURU	784	1	1	4	3
15	GADIMUDIDAM	897	10	2	0	0
16	KANCHARAM1	859	2	2	0	0
17	KANCHARAM2	823	2	1	0	0
18	LATCHAYYETA-01	1606	27	6	1	0
19	KONDAMPETA	915	37	10	0	0
20	MITTIREDDYSTREET	926	8	4	0	0
21	MADIGAVEEDHI	1377	77	29	5	0
22	INDIRAMMACOLONY	903	3	2	1	0
23	BUTCHIMPETA	973	1	1	1	0
24	PONUGUTIVALASA	1366	17	10	1	11
25	ADARSHNAGAR	1047	9	2	4	15
26	KOTHAVALASA	1005	14	6	1	0
27	SATYANARAYANAPURAM	828	11	5	0	0
28	LATCHAYYETA-02	868	6	4	15	15

and the Choi et al. [9] proposed HerDing which is a herb recommended system useful for treating a chronic disease like leukaemia, diabetics, heart diseases, etc. The HerDing recommends respective natural products (herbs) based on symptoms of a patient.

In the literature, recommended systems [5–9] using different machine learning techniques were proposed. In addition to this, recently, Srinivasa Rao et al. [10] proposed a machine learning-based web survey through mobile phone for the improvement in the identification of possible COVID-19 cases and to reduce the spread of the virus among susceptible people who are under quarantine. The survey intention is to collect travel history and symptoms from a person who is under COVID-19 investigation. Similarly, AI techniques are applied on such preliminary data for early identification of people with COVID-19 and non-COVID-19 symptoms as well as to classify them into no-risk, minimal-risk, moderate-risk, and high-risk groups. The identified high-risk people can be recommended for isolation. In line with the proposed web survey through mobile phone [10], in this paper, we proposed COVID-19 test recommendation system using machine learning techniques on

house-to-house survey data collected by the health department of Andhra Pradesh (AP) state government to identify the people infected with General and High-risk symptoms. This proposed work recommends people for COVID-19 test based on disease class – No-Symptoms, General Symptoms, Low-risk, Medium-risk, and High-risk – identified using machine learning methods like KNN, DT, Naïve Bayes, RF, SGD, and SVM. The proposed symptom-based test recommendation system gives more than 99% accuracy in recommending COVID-19 test to the people who suffer from defined ill-health symptoms described in Algorithms 1 and 2. These algorithms are prepared based on inputs of the Superintendent of Physical Health Center (Rajam) mentioned in Table 2 and representation of each symptom is as shown in Fig. 1.

In this piece of work, the proposed Algorithms 1 and 2 are applied on house-to-house surveyed data to build a synthesized data set and then, data balancing methods applied to this synthesized data set to make it balanced. Now, the machine learning techniques like KNN, DT, Naïve Bayes, RF, SGD, and SVM are applied on a both synthesized and balanced data set to identify different

Table 2  
Description of data set shown in Table 1

Secretariat	Is an government office at each village.
Number-of-Houses	The total number of houses under each Secretariat.
General Public Symptomatic	The number of people suffering from general symptoms – Cold, Cough and Fever problem – in each Secretariat.
High Risk Factors	<p>Divided into three categories by considering three symptoms such as Age, Breathing and Comorbidites along with general symptoms.</p> <p>(1) People who are above sixty years, have Comorbidities, suffering from any of the general symptom and Breathing problem considered as <b>High Level</b>.</p> <p>(2) People who are below sixty years, have Comorbidities, having general symptoms and breathing problem considered as <b>Medium Level</b>.</p> <p>(3) People with above sixty years with general symptoms considered as <b>Low Level</b>.</p> <p><b>Note:</b> that the chronic diseases like blood pressure, diabetics, leukemia, and etc. are termed as <b>Comorbidities</b>.</p>
General Symptomatic and Age > 60	The number of people who are above 60 years and having general symptoms in each Secretariat.
Age > 60 and comorbidities	The number of people who are above 60 years, having Comorbidities and having general symptoms in each Secretariat.
Age <= 60 and comorbidities	The number of people who are below 60 years, having Comorbidities and having general symptoms in each Secretariat.

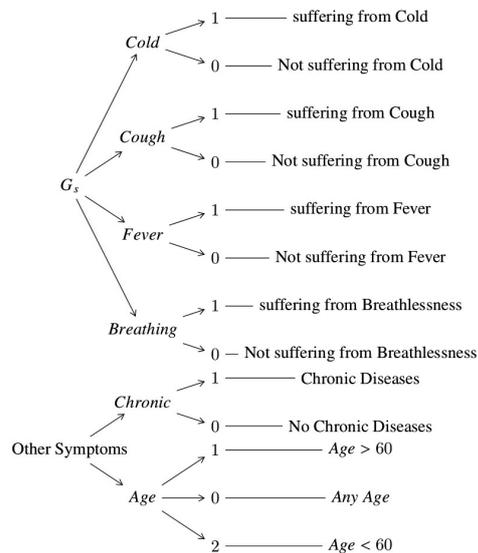


Fig. 1. The Symptom and it's representation.

classes based on age, comorbidity, and other symptoms (which are defined by the health department of AP) of people suffering from various diseases.

### 3. Symptom based COVID-19 test recommendation system using machine learning techniques

The recent statistics reveal that spread of the COVID-19 virus transformed into a pandemic form epidemic. The mankind of the entire world is fighting against the COVID-19 pandemic. The World Health Organization (WHO) [11,12] strictly recommends more tests to ad-

mit COVID-19 infected people in COVID-Hospitals to decrease community spread of COVID-19. It also recommends different testing priorities and strategies [12] for early detection of infected people as well as best utilization of available medical – testing infrastructure, medical staff, etc. – resources. The proposed COVID-19 testing recommender system is very useful to detect and separate probable high-risk category people. The flow of proposed work is as shown in Fig. 2.

#### 3.1. About data set

To address the COVID-19 testing problem, the Andhra Pradesh (AP) state government of India conducted a house-to-house survey in the entire state through village/Town level employees to find the health status of people. Because of this, the data set consisting of a rural and urban population of Rajam Mandal is collected from the Primary Health Center (PHC) of Rajam, Srikakulam district. Table 1 shows the Secretariat (an AP government office established for one village or group of villages based on population) wise survey data. It consists of fields like the name of Secretariat, the number of houses visited under that Secretariat, public suffering from General symptoms, and three levels of risk factors  $R_f$ . The Table 2 consists detailed explanation of each feature present in the data set shown in Table 1. For example, under BODDAM Secretariat 1646 houses were visited in which 12 persons are suffering from any of the general symptom (cold, cough, and fever), 6 persons are above sixty years of age as well as suffering from any of the general symptoms, 11 persons who have above sixty years are suffering from

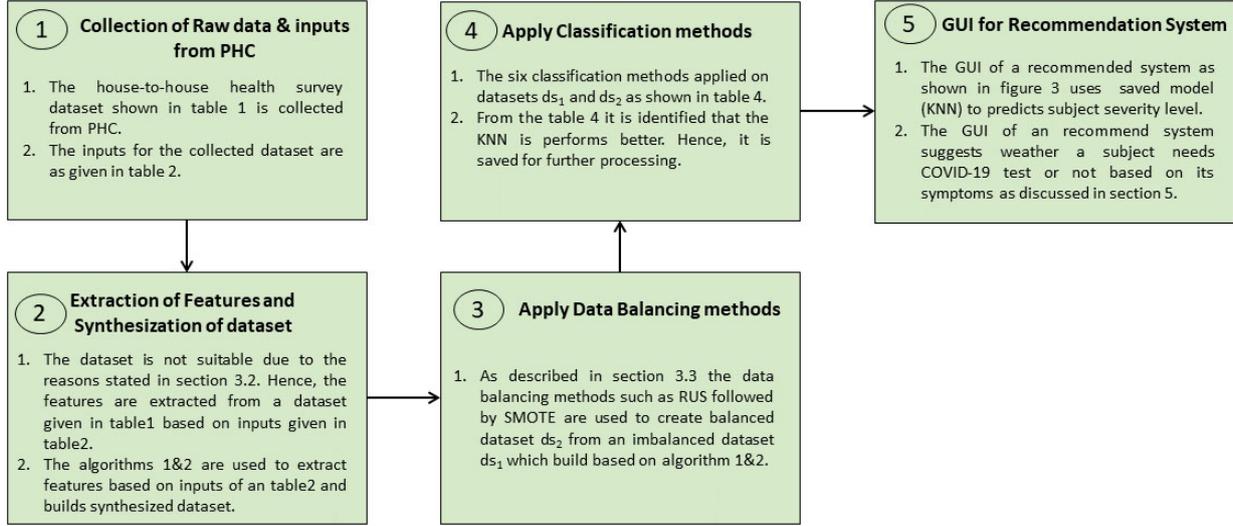


Fig. 2. The flow of proposed work.

both chronic diseases and any of the general symptom and finally, 11 people who are less than or equal to sixty years suffering from comorbidities and general symptoms.

**Algorithm 1:** Synthesized data set creation by initializing all the features

**Purpose:** Initialization of features such as Cold, Cough, Fever, Breath, Age, Chronic and Recommendation.

**Description:** The variables such as  $Cold_{1 \times T_p}$ ,  $Cough_{1 \times T_p}$ ,  $Fever_{1 \times T_p}$ ,  $Breath_{1 \times T_p}$ ,  $Age_{1 \times T_p}$ ,  $Chronic_{1 \times T_p}$ , and  $Recommendation_{1 \times T_p}$  indicated as Cold, Cough, Fever, Breath, Age, Chronic and Recommendation respectively. Each feature is an one dimensional array of size  $1 \times T_p$ .

**Input:** The extracted features as in step 3 of Section 3.2 considered as input.

```

for i ← 0 to  $T_p$  by 1 do
  Cold[i] ← 0;
  Cough[i] ← 0;
  Fever[i] ← 0;
  Breath[i] ← 0;
  Age[i] ← 0;
  Chronic[i] ← 0;
  Recommendation[i] ← 0;
end

```

**end**

call **set\_gs\_rf** ( $\theta$ ,  $I$ ,  $N_s$ )

call **set\_gs\_rf** ( $I$ ,  $N_s + 1$ ,  $g_s$ )

call **set\_gs\_rf** ( $2$ ,  $g_s + 1$ ,  $r_l$ )

call **set\_gs\_rf** ( $3$ ,  $r_l + 1$ ,  $r_m$ )

call **set\_gs\_rf** ( $4$ ,  $r_m + 1$ ,  $r_h$ )

**Output:** Make each feature  $Cold_{1 \times T_p}$ ,  $Cough_{1 \times T_p}$ ,  $Fever_{1 \times T_p}$ ,  $Breath_{1 \times T_p}$ ,  $Age_{1 \times T_p}$ ,  $Chronic_{1 \times T_p}$  and  $Recommendation_{1 \times T_p}$  as individual columns to the data set  $d_{s1}$ .

### 3.2. Implementation of rules to create synthesized data set

The original data set shown in Table 1 is imbalanced and not suitable for the learning process because of the below reasons

- Huge set of people without symptoms in the given data set.
- The number of people in each house is not known since it populates secretariat wise data.
- The populated data shows how many people under one secretariat have symptoms in each category such as the total number of persons either male or female who has general symptoms ( $G_s$ ) and High-Risk Factor (described in Table 2).
- The symptoms – cold, cough, fever, or breathing – of a patient is unknown in each category.

In this connection, the below steps are implemented to extract features based on the data given in Tables 1 and 2 to address the above-said reasons.

**Step 1:** According to the Indian context, it is assumed that the total number of people who dwell in each house is four. So, the number of houses multiplied by four is the total population in each secretariat. As a result, the total surveyed population  $S_p$  is the sum of Secretariat wise houses multiplied by four. That is  $28326 \times 4 = 113304$ .

**Step 2:** According to the inputs of PHC superintendent a subject does not have any symptoms treated as No-Symptoms ( $N_s$ ), subject with

Cough, Cold, Fever are classified as general symptoms ( $G_s$ ), the Age and Comorbidities (chronic diseases) are considered as two other important features along with  $G_s$  to decide risk factors ( $R_f$ ) such as Low-risk ( $r_l$ ), Medium-risk ( $r_m$ ), and High-risk ( $r_h$ ). The subject who have  $age > 60$  with  $G_s$  and breathing problems considered as  $r_l$ , the subject who have  $age < 60$  with  $G_s$ , breathing and Comorbidities considered as  $r_m$ , and the subject who have  $age > 60$  with  $G_s$ , breathing and Comorbidities considered as  $r_h$ .

- Step 3: The features or symptoms such as Cold, Cough, Fever, Breathing, Chronic (Comorbidities), and Age are identified according to step 2
- Step 4: The value 1 (one) of a corresponding feature represents that the people suffering from that symptom and value 0 (zero) means not suffering from the corresponding symptom as shown in Fig. 1.
- Step 5: Among  $S_p$ , 321 people suffering from general symptoms, 123 are classified as a Low-risk category, 67 are in Medium-risk and 77 are in the High-risk category.
- Step 6: Based on the data set shown in Table 1 the symptoms of a person who is under  $G_s$ , Low-risk, Medium-risk, and High-risk category is unknown. In this context, it can be assumed that the person who may suffer from  $G_s$  may suffer from either only cold, only cough, only fever, only cold and cough, only cold and fever, only cough and fever, or all three symptoms. Hence, each person who suffers from  $G_s$  multiplied by 7 to cover all possibilities.
- Step 7: In the same way as in step 6 the people who suffer from  $R_f$  are multiplied by 16 since  $R_f$  considers 6 features such as cold, cough, fever, breathing, age, and chronic. Among these symptoms the age and chronic are fixed with either 0 (zero) or 1 (one) based on the category of risk factor  $R_f$  and the remaining four features keep changing.
- Step 8: As per step 6&7 multiply the total people under  $G_s$  category with 7, Low-risk (123) category with 16, Medium-risk (77) category with 16 and High-risk (67) category with 16. So, the  $S_p$  is added with  $321 \times 7$  ( $g_s$ ),  $123 \times 16$  ( $r_l$ ),  $77 \times 16$  ( $r_m$ ) and  $67 \times 16$  ( $r_h$ ).

Hence, the total size of each feature is  $1 \times T_p$  where  $T_p = S_p + g_s + r f_l + r f_m + r f_h$ . The remaining 113192 entries are considered as No-Symptoms.

These steps are used to synthesize the data set according to the data given in Table 1 by extracting the necessary features based on inputs of PHC superintendent (Rajam) given in Table 2 followed by providing proper initialization of extracted features to make them

---

**Algorithm 2:** Initialization of all possibilities according to step 6 and 7 as in Section 3.2

---

**Purpose:** To initialize symptoms under each class for all possible cases.

**Input:** The class number(0, 1, 2, 3, and 4), starting index and end index of corresponding class considered as input

**Description:** Each feature is filled with either 1 or 0 to represent all five classes as described in step 8 of Section 3.2.

**set\_gs\_rf (class, start, end)**

Initialize array  $a[] \leftarrow \{0, 0, 0, 0\}$

Initialize counter  $c \leftarrow 0$

**for**  $i \leftarrow start$  **to**  $end$  **do**

**if**  $c == 0$  **then**

$a[0] \leftarrow 0$ ;  $a[1] \leftarrow 0$ ;  $a[2] \leftarrow 0$ ;  $a[3] \leftarrow 0$ ;

**else**

**for**  $j \leftarrow 0$ ;  $c > 0$ ;  $j + +$  **do**

$a[j] \leftarrow c \% 2$ ;

$c \leftarrow c / 2$ ;

**end**

**end**

  Cold[i]=a[0]; Cough[i]=a[1]; Fever[i]=a[2];

**if**  $class == 0$  **then**

    Recommendation[i]=0 (No-Symptoms);

**end**

**if**  $class == 1$  **then**

    Breath[i]=a[3]; Age[i]=0; Chronic[i]=0;

    Recommendation[i]=1 ( $G_s$ );

**end**

**if**  $class == 2$  **then**

    Breath[i]=a[3]; Age[i]=2; Chronic[i]=1;

    Recommendation[i]=2 ( $r_l$ );

**end**

**if**  $class == 3$  **then**

    Breath[i]=a[3]; Age[i]=1; Chronic[i]=0;

    Recommendation[i]=3 ( $r_m$ );

**end**

**if**  $class == 4$  **then**

    Breath[i]=a[3]; Age[i]=1; Chronic[i]=1;

    Recommendation[i]=4 ( $r_h$ );

**end**

$c++$

**if**  $class == 0$  and  $c > 7$  **then**

$c = 0$

**end**

**if**  $class \neq 0$  and  $c > 15$  **then**

$c = 0$

**end**

**end**

**Output:** An index of each feature consists a value either 0 or 1 to indicate corresponding symptom.

---

Table 3  
The class labels of data sets  $d_{s_1}$  and  $d_{s_2}$  before and after train-test split

Classes	$d_{s_1}$	$d_{s_2}$		After splitting of $d_{s_1}$		After splitting of $d_{s_2}$	
		RUS	SMOTE	Train (70%)	Test (30%)	Train (70%)	Test (30%)
0	113192	1072	2247	79200	33992	749	323
1	2247	2247	2247	1572	675	1590	657
2	1968	1968	2247	1397	571	1359	609
3	1072	1072	2247	757	315	764	308
4	1232	1232	2247	871	361	851	381

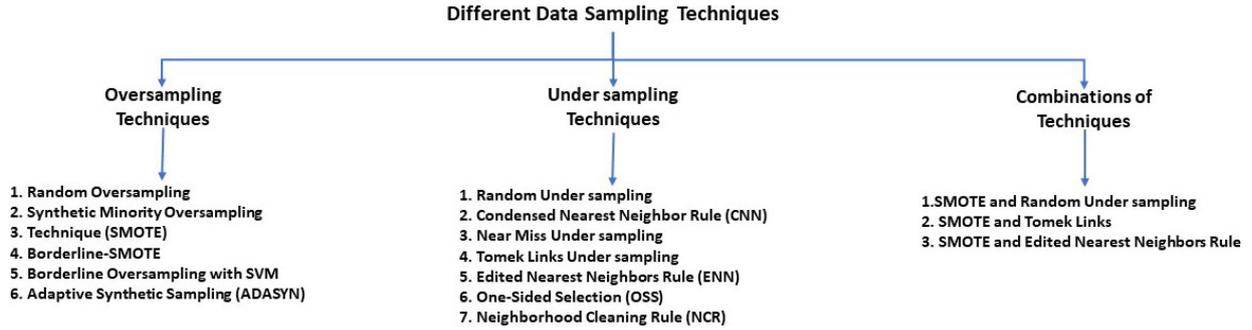


Fig. 3. The data balancing methods

suitable for classification as described in Algorithms 1 and 2. Finally, prepare the data set  $d_{s_1}$  by combining all initialized features. Based on a literature study it is observed that the training and prediction accuracy of the proposed model is effectively evaluated with a balanced data set. Hence, the data balancing techniques discussed in Section 3.3 are applied on a resultant data set  $d_{s_1}$  to generate even more balanced data set  $d_{s_2}$ . The different machine learning methods for classification applied on both data sets ( $d_{s_1}$  &  $d_{s_2}$ ) and its results are analyzed in Section 4.

### 3.3. Data balancing techniques

The data re-sampling methods are widely used techniques to make imbalanced class distribution of a respective data set as balanced distribution. The approaches such as under-sampling and over-sampling are two variants for data re-sampling. The uneven train and test data split of class labels of an imbalanced data set is a major drawback. The data sampling techniques shown in Fig. 3 are used to make an equal number of class labels of a respective data set through either oversampling or under-sampling methods. In this piece of work, RUS followed by SMOTE [13–15] is used to create a balanced data set  $d_{s_2}$ . Initially, it randomly deletes some example entries from majority class (0<sup>th</sup> → No-Symptoms) of  $d_{s_1}$  and makes it equal to the minority class (3<sup>rd</sup> → Medium-Risk) through random

under-sampling technique. Secondly, the oversampling of minority classes is performed through SMOTE to make all classes with an equal number of examples. The entries of each class label for both the data sets –  $d_{s_1}$  and  $d_{s_2}$  – is shown in Table 3.

### 3.4. Classification techniques

The supervised learning method [16,17] like classification has two variants such as Binary and Multi-class classifiers are used to classify entries of a data set into the identified class labels. A binary classifier classifies a subject into either of two classes 0 or 1, Yes or No, Male or Female, True or False, etc. A Multiclass classifier works on a data set with more than two class labels and classifies a subject into either of those classes.

The classification methods such as KNN, DT, Naive Bayes, RF, SGD and SVM are applied on both  $d_{s_1}$  and  $d_{s_2}$ . The prediction time is high in KNN [18,19] and training time is very less since it only stores training data and applies logic when it receives testing data for prediction. Hence, it is called as a lazy learner. The value of  $K$  is the most important parameter in K-NN. This algorithm performs three steps during the prediction process. Firstly, the distance between a new data point and all data points of the data set is measured. Secondly, finds  $K$  number of data points from a data set that has less distance from the new data point. Finally, the class label of the majority of the points among  $K$

number of data points is selected as the class of the new data point.

The DT [19,20] takes more time for training and less time for prediction and it builds a classification model during its training process. It is one of the popular supervised learning techniques used to solve both classification and regression problems, but it is widely used to classify given data set for accurate decision making. The DT builds a tree for a given data set which consists of two nodes such as Decision-node (Root-node) and Leaf-node. The starting point of the tree is called as Root-node. The Decision-node consists of multiple branches each branch shows a direction towards a final decision. The Leaf-node does not contain branches and gives the final result for the given conditions. The Naïve Bayes classification model is most suitable for large data sets, it is quick in prediction, performs well for multi-class prediction, and needs less training data. This model is designed based on the Bayes theorem. RF is a less biased and less data-sensitive ensemble learning model. It creates a specified number of DT instances on a given data set and takes the majority opinion of DT instances as an outcome.

SGD [21] is one variant of gradient descent optimization algorithms. It takes a very small amount of data from a training set for each iteration to compute the gradient and updates its weight matrix subsequently for each iteration to quickly reach an optimal point. Hence, this algorithm is much faster than other gradient descent algorithms and very useful for huge training sets for quick converge.

SVM is a supervised learning algorithm used for both classification and regression problems. However, it is widely used in classification. It finds a right hyper-plane that differentiates all classes accurately.

#### 4. Evaluation of synthesized data set and classification models

The performance of machine learning models is evaluated using accuracy, sensitivity, specificity, precision, rate of misclassification, and F1-score metrics [19]. The calculation of evaluation metrics uses numerical values of true positive (TP), true negative (TN), false positive (FP), and false-negative (FN) which are obtained from confusion matrices of a respective classification model is as shown below:

- Accuracy =  $(TP + TN)/(TP + TN + FP + FN)$
- Sensitivity or Recall or true positive rate =  $TP/(TP + FN)$

- Specificity or True negative rate =  $TN/(TN + FP)$
- Precision =  $TP/(TP + FP)$
- Rate of mis-classification =  $(FP + FN)/(TP + TN + FP + FN)$
- F1-measure =  $2 * (\text{precision} * \text{recall})/(\text{precision} + \text{recall})$

Four steps are performed in this piece of work. At first, the COVID-19 survey data shown in Table 1 is not suitable for learning as described in Section 3.2. Hence, the Algorithms 1 and 2 are applied on Table 1 to create synthesized data set  $d_{s_1}$ . Second, the data balancing methods – described in Section 3.3 – such as random under-sampling followed by SOMTE data balancing methods applied on  $d_{s_1}$  to create balanced data set  $d_{s_2}$ . Third, six classification models such as KNN, DT, Naïve Bayes, RF, SGD and SVM are applied on both the data sets  $d_{s_1}$  and  $d_{s_2}$ . Finally, The less data sensitive and data biased KNN learning model is integrated with the GUI of the COVID-19 Test Recommender system.

In first task, the data set  $d_{s_1}$  is created as described in Algorithm 1. According to the step-1&8 of Algorithm 1 the total number of houses is 28326 based on data shown in Table 2 multiplied with four to fix the  $S_p$ . Then, the people of  $g_s, r_l, r_m,$  and  $r_h$  added to  $S_p$ . As a result, the  $T_p$  in  $d_{s_1}$  is 119823. The data samples of each class of  $d_{s_1}$  shown in Table 3.

In the second task, the RUS data balancing method makes data samples of a maximum category is equal to the minority class of the respective data set by deleting the required number of entries from the maximum category. On the other hand, the data oversampling method SMOTE makes the number of data samples of all classes is equal by adding the required number of extra data samples to each class which are less than the maximum data entries class of the same data set. The RUS followed by SMOTE applied on  $d_{s_1}$  to create  $d_{s_2}$ . In this process, the RUS reduces data entries (113192) of a maximum category (No Symptoms) to data entries (1072) of the minority category (Medium-risk). As a result, 1072 data entries in No-Symptoms category, 2247 samples in General symptoms, 1968 data entries in  $r_l$ , 1072 data samples in  $r_m$  and 1232 entries in  $r_h$  as shown in Table 3. Next, the minority data oversampling method SMOTE is applied to the result of RUS to make an equal number of example data samples to all five categories. The result of SMOTE is stored in  $d_{s_2}$  which is shown in Table 3.

In the third task, The training and testing accuracy of classification models like KNN, DT, Naïve Bayes, RF, SGD, and SVM is applied on both  $d_{s_1}$  and  $d_{s_2}$  with 70:30 percent of train and test ratio. Then, the

Table 4  
The training and testing accuracy of all six classification methods on  $d_{s_1}$  and  $d_{s_2}$

Classification methods	$d_{s_1}$		$d_{s_2}$		$d_{s_1}$		$d_{s_2}$	
	Training accuracy	Testing accuracy	Training accuracy	Testing accuracy	Training time	Testing time	Training time	Testing time
K-nearest neighbor	1.0	1.0	1.0	1.0	1.31	40.25	0.02	0.06
Decision tree	0.99	1.0	0.99	1.0	0.06	0.01	0.003	0.001
Naive bayes	0.99	1.0	0.99	1.0	0.08	0.01	0.003	0.002
Random forest	0.99	1.0	0.99	1.0	1.13	0.39	0.19	0.03
Stochastic gradient descent	0.99	1.0	0.99	1.0	0.49	0.21	0.015	0.004
Support vector machine	0.99	1.0	0.99	1.0	2.39	0.76	0.25	0.07

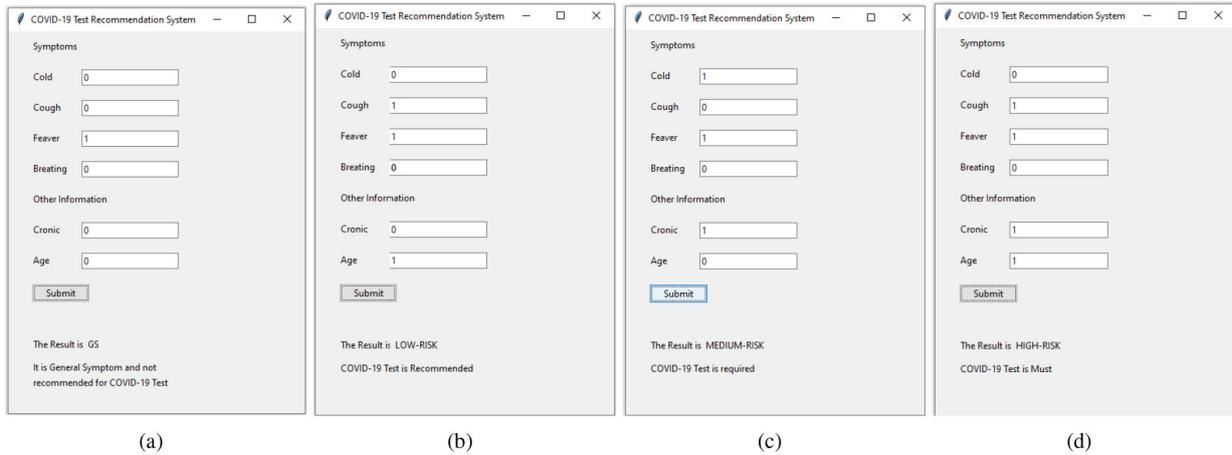


Fig. 4. (a) General symptom, (b) Low-risk, (c) Medium-risk, (d) High-risk.

accuracy and response times of training and testing of each classification model on both the data sets are shown in Table 4. Moreover, the evaluation metrics of all these classification models on both  $d_{s_1}$  and  $d_{s_2}$  are also evaluated resulting in sensitivity or true positive rate is equal to 1.0, the specificity or true negative rate is equal to 0, precision is 1.0, rate of misclassification is 0, and F1-score is 1.0.

From the results as shown in Table 4 KNN classifier performs well on both the datasets when compared with other classifiers. It is a multi-class classifier suits to the datasets  $d_{s_1}$  and  $d_{s_2}$  which consists five classes, consumes less time for both training and prediction, and it performs well when the size of a dataset is small as well as training data is larger than the number of features. Hence, it is saved and integrated with the COVID-19 Recommended system as shown in Fig. 4 as the final task.

## 5. The symptom based recommended system

The results discussed in Section 4 shows the performance of all six classification methods such as KNN,

DT, Naïve Bayes, RF, SGD, and SVM gives better results on both the data sets. However, the KNN outperformed other classifiers with both training and testing data. Henceforth, it is saved and used in the GUI of a recommended system to predict the status of a person based on collected symptoms. The proposed recommended system collects symptoms of Cold, Cough, Fever, Breath, Chronic (Comorbidities), and Age from the user as shown in Fig. 4. This information is used to recommend whether the COVID-19 test is required or not as well as the level of severity in collected symptoms. The proposed recommended system is useful for both web and mobile-based survey. The symptoms of a subject are entered through the GUI shown in Fig. 4. A health assistant can enter either 0 or 1 in a specific text field of the GUI. As shown in Fig. 1 the value 0 (Zero) indicates the no presence of a symptom, and the value 1 (One) indicates the presence of a symptom.

The machine learning algorithms are applied on the synthesized dataset where the algorithms are adopted from the sklearn API. All the modules in the proposed work such as creation of synthesized data, data balancing, classification, saving of model and GUI are implemented in python. The source code of this work is avail-

able at <https://github.com/KLakshmanarao/covid19> for reference.

## 6. Conclusion and future work

We have evaluated our proposed model with six classification algorithms. From the experimentation results, it is observed that the evaluation metrics – Sensitivity, Specificity, Precision, Rate of misclassification, and F1-Score – and train-test accuracy are good and similar on both the data sets  $d_{s_1}$  and  $d_{s_2}$ . Consequently, it is proved that the generation of synthesized data set  $d_{s_1}$  through Algorithms 1 and 2 is balanced. It is also observed that the integration of the COVID-19 test recommender system with the KNN model achieves 100% accuracy in predicting the category of a subject based on its symptoms. In addition to this, the proposed COVID-19 Test recommended system using machine learning techniques will be very helpful to public health systems to decide whether a subject needs a COVID-19 test or not. In this proposed model, all the classes are recommended for the COVID-19 test except for the subject with general symptoms. Because of the scarcity of COVID-19 test kits and other medical resources, the proposed recommender system will be very useful for optimal utilization of medical resources since only required people will be recommended for COVID-19 testing. If a subject which is recommended for COVID-19 test can be quarantined to reduce the spread of the virus. The citizen can also use the proposed system and test themselves on their own. The proposed work can be extended in such a manner that the integration of additional symptoms related to COVID-19 can be taken into consideration to make the prediction more active and robust.

## References

- [1] Krishnakumar B, Rana S. COVID 19 in INDIA: Strategies to combat from combination threat of life and livelihood. *Journal of Microbiology, Immunology and Infection*. 2020 Jun 1; 53(3): 389–91.
- [2] Barstugan M, Ozkaya U, Ozturk S. Coronavirus (covid-19) classification using ct images by machine learning methods. *arXiv preprint arXiv:2003.09424*. 2020 Mar 20.
- [3] Yang Z, Zeng Z, Wang K, Wong SS, Liang W, Zanin M, Liu P, Cao X, Gao Z, Mai Z, Liang J. Modified SEIR and AI prediction of the epidemics trend of COVID-19 in China under public health interventions. *Journal of Thoracic Disease*. 2020 Mar; 12(3): 165.
- [4] Tanabe K. Society coexisting with COVID-19. *Infection Control & Hospital Epidemiology*. 2020 Aug; 41(8): 988–9.
- [5] Carter JA, Long CS, Smith BP, Smith TL, Donati GL. Combining elemental analysis of toenails and machine learning techniques as a non-invasive diagnostic tool for the robust classification of type-2 diabetes. *Expert Systems with Applications*. 2019 Jan 1; 115: 245–55.
- [6] Ali M, Thanh ND, Van Minh N. A neutrosophic recommender system for medical diagnosis based on algebraic neutrosophic measures. *Applied Soft Computing*. 2018 Oct 1; 71: 1054–71.
- [7] Chen RC, Huang YH, Bau CT, Chen SM. A recommendation system based on domain ontology and SWRL for anti-diabetic drugs selection. *Expert Systems with Applications*. 2012 Mar 1; 39(4): 3995–4006.
- [8] Phanich M, Pholkul P, Phimoltares S. Food recommendation system using clustering analysis for diabetic patients. In 2010 International Conference on Information Science and Applications. IEEE. 2010 Apr 21. pp. 1–8.
- [9] Choi W, Choi CH, Kim YR, Kim SJ, Na CS, Lee H. HerDing: herb recommendation system to treat diseases using genes and chemicals. *Database*. 2016 Jan 1; 2016.
- [10] Rao AS, Vazquez JA. Identification of COVID-19 can be quicker through artificial intelligence framework using a mobile phone-based survey when cities and towns are under quarantine. *Infection Control & Hospital Epidemiology*. 2020 Jul; 41(7): 826–30.
- [11] Al-Muharraqi MA. Testing recommendation for COVID-19 (SARS-CoV-2) in patients planned for surgery-continuing the service and ‘suppressing’ the pandemic. *The British Journal of Oral & Maxillofacial Surgery*. 2020 Jun; 58(5): 503.
- [12] World Health Organization. Laboratory testing strategy recommendations for COVID-19: interim guidance, 21 March 2020. World Health Organization; 2020.
- [13] Pecorelli F, Di Nucci D, De Roover C, De Lucia A. A large empirical assessment of the role of data balancing in machine-learning-based code smell detection. *Journal of Systems and Software*. 2020 Nov 1; 169: 110693.
- [14] Susan S, Kumar A. SSOMaj-SMOTe-SSOMin: Three-step intelligent pruning of majority and minority samples for learning from imbalanced datasets. *Applied Soft Computing*. 2019 May 1; 78: 141–9.
- [15] Alghamdi M, Al-Mallah M, Keteyian S, Brawner C, Ehrman J, Sakr S. Predicting diabetes mellitus using SMOTE and ensemble machine learning approach: The Henry Ford Exercise Testing (FIT) project. *PloS One*. 2017 Jul 24; 12(7): e0179805.
- [16] Singh A, Thakur N, Sharma A. A review of supervised machine learning algorithms. In 2016 3rd International Conference on Computing for Sustainable Global Development (INDIACom). IEEE. 2016 Mar 16. pp. 1310–1315.
- [17] Alpaydin E. Introduction to machine learning. MIT press; 2020 Mar 24.
- [18] Friedman JH, Baskett F, Shustek LJ. An algorithm for finding nearest neighbors. *IEEE Transactions on Computers*. 1975 Oct; 100(10): 1000–6.
- [19] Choudhury A, Gupta D. A survey on medical diagnosis of diabetes using machine learning techniques. In *Recent Developments in Machine Learning and Data Analytics*. Springer, Singapore. 2019. pp. 67–78.
- [20] Argentiero P, Chin R, Beaudet P. An automated approach to the design of decision tree classifiers. *IEEE Transactions on Pattern Analysis and Machine Intelligence*. 1982 Jan; 4(1): 51–7.
- [21] Ruder S. An overview of gradient descent optimization algorithms. *arXiv preprint arXiv:1609.04747*. 2016 Sep 15.
- [22] Bernheim A, Mei X, Huang M, Yang Y, Fayad ZA, Zhang N, Diao K, Lin B, Zhu X, Li K, Li S. Chest CT findings in

- coronavirus disease-19 (COVID-19): relationship to duration of infection. *Radiology*. 2020 Feb 20: 200463.
- [23] Ai T, Yang Z, Hou H, Zhan C, Chen C, Lv W, Tao Q, Sun Z, Xia L. Correlation of chest CT and RT-PCR testing for coronavirus disease 2019 (COVID-19) in China: A report of 1014 cases. *Radiology*. 2020 Aug; 296(2): E32–40.
- [24] Shi H, Han X, Jiang N, Cao Y, Alwalid O, Gu J, Fan Y, Zheng C. Radiological findings from 81 patients with COVID-19 pneumonia in Wuhan, China: A descriptive study. *The Lancet Infectious Diseases*. 2020 Apr 1; 20(4): 425–34.
- [25] Hanke H, Knees D. A phase-field damage model based on evolving microstructure. *Asymptotic Analysis*. 2017 Jan 1; 101(3): 149–80.
- [26] Lefever E. A hybrid approach to domain-independent taxonomy learning. *Applied Ontology*. 2016 Jan 1; 11(3): 255–78.
- [27] Meltzer PS, Kallioniemi A, Trent JM. Chromosome alterations in human solid tumors. *The genetic basis of human cancer*. New York: McGraw-Hill. 2002: 93–113.
- [28] Murray PR, Rosenthal KS, Pfaller MA. *Medical microbiology* E-book. Elsevier Health Sciences; 2020 Mar 10.
- [29] Wilson E. Active vibration analysis of thin-walled beams (Doctoral dissertation, Ph. D. Dissertation, University of Virginia).