

Application of Machine Learning K-Means Clustering and Linear Regression in Determining the Risk Level of Pulmonary Tuberculosis

1st Abhijit Pathak

Assistant Professor

BGC Trust University Bangladesh

Chattogram, Bangladesh

abhijitpathak@bgctub.ac.bd

0000-0001-7734-0271

2nd Ziaul Islam Bablu

Research Assistant

BGC Trust University Bangladesh

Chattogram, Bangladesh

ziaulislambablu2@gmail.com

0009-0000-8783-8022

3rd Towhidul Haque Limon

Research Assistant

BGC Trust University Bangladesh

Chattogram, Bangladesh

towhidulhaque4455@gmail.com

0009-0002-4177-6991

4th Sowmik Barua

Research Assistant

BGC Trust University Bangladesh

Chattogram, Bangladesh

sowmikbarua7878@gmail.com

0009-0004-4974-3478

5th Piyal Dey

Research Assistant

BGC Trust University Bangladesh

Chattogram, Bangladesh

piyaldey6@gmail.com

0009-0000-8385-1617

6th Mowmita Tajnin Jiba

Research Assistant

BGC Trust University Bangladesh

Chattogram, Bangladesh

mowmitatajninj@gmail.com

0009-0000-7578-4813

7th Touhidul Alam Seyam

Research Assistant

BGC Trust University Bangladesh

Chattogram, Bangladesh

touhidulalam@bgctub.ac.bd

0009-0007-7512-1893

Abstract—Pulmonary tuberculosis (TB) remains a significant public health concern in densely populated regions like Bireuen, Bangladesh, which reported 755 cases in 2019 among a population of 400,000. This study used data from Bangabandhu Sheikh Mujib Medical University Hospital and the Health Department across 17 districts to identify high-risk areas and predict disease incidence. Utilizing K-Means clustering and Cluster-wise Regression, the analysis identified two high-risk areas in Cluster 1, six in Cluster 2, and nine in Cluster 3, with a regression analysis R-squared value of 0.5740, indicating moderate predictive capacity. These findings provide critical insights for public health authorities to devise targeted interventions and allocate resources effectively. Strategies such as targeted screening programs and improved access to diagnostic and treatment facilities in high-risk areas can help mitigate TB's impact. The study emphasizes the importance of continued surveillance, monitoring, and collaborative efforts among government agencies, healthcare providers, researchers, and community stakeholders to achieve TB elimination in Bangladesh.

Index Terms—Pulmonary Tuberculosis, Linear Regression, K-Means Clustering, Predictive modeling, Disease mapping.

I. INTRODUCTION

Pulmonary tuberculosis (TB) is a major global health challenge, with India having the highest number of cases. According to the World Health Organization (WHO), TB is one of the top 10 diseases causing death globally. This study aims to enhance TB risk assessment using machine learning

techniques like K-Means clustering and linear regression. By analyzing datasets on population density, geographical distribution, and TB patient numbers, the authors identify high-risk areas and forecast disease trends. K-Means clustering groups districts based on TB distribution patterns, while cluster-wise linear regression improves prediction accuracy by considering variable interrelationships. The authors employ these techniques to identify distinct patterns and assess the relationship between specific TB risk factors and the likelihood of developing pulmonary TB. Evaluation against traditional methods will gauge predictive capabilities. This comparative analysis will help determine the added value of machine learning approaches in TB risk prediction. The insights derived from the machine learning models will highlight the most influential risk factors contributing to the development of pulmonary TB. By analyzing these factors within each identified cluster, the authors can provide a detailed understanding of how different variables interplay to affect TB risk. This information is crucial for developing targeted interventions. Based on the identified risk factors and clusters, the authors will suggest potential interventions or strategies for TB prevention and control. These may include targeted public health campaigns, personalized medical monitoring, and specific lifestyle or environmental modifications. The goal is to utilize the insights gained from the analysis to inform

effective and targeted TB prevention and control measures, ultimately reducing the incidence and impact of pulmonary tuberculosis. In Bangladesh, efforts to combat tuberculosis are ongoing through programs like the NTP. However, the risk level of pulmonary tuberculosis remains a concern due to undetected cases and challenges in providing effective treatment and prevention measures. By utilizing innovative machine learning approaches, the authors aim to improve TB risk assessment and stratification, leading to better-targeted interventions and control strategies. The application of these advanced techniques can potentially revolutionize the way TB risk is evaluated and managed, offering a scalable and precise method to address this persistent global health issue.



Fig. 1. Stages to determine the Risk Level of Pulmonary Tuberculosis.

In this study, the authors aim to use advanced machine learning techniques to understand pulmonary tuberculosis (TB) epidemiology and guide prevention and control strategies. They plan to:

- Implement K-Means clustering to identify TB risk patterns.
- Stratify individuals into risk groups based on demographics and clinical data.
- Apply linear regression to assess TB risk factors within each cluster.
- Evaluate model performance against traditional methods.
- Identify influential TB risk factors.
- Suggest targeted interventions for TB prevention and control.

II. RELATED WORKS

The reviewed paper provides a thorough examination of pulmonary tuberculosis (TB), detailing the lung as the primary infection site, symptoms, laboratory examinations, chest radiography, and CT scans. It discusses TB in the elderly, those on anti-tumor necrosis factor alpha inhibitors, and the diagnosis of pleural TB. The paper also covers pulmonary TB in HIV patients and its complications [1]. A significant portion addresses childhood pulmonary TB, highlighting three key concepts from chemotherapy literature: accurate case definitions, risk stratification, and the diverse spectrum of disease pathology. These concepts are crucial for diagnosing and treating childhood TB [2]. The paper compares risk factors for extra-pulmonary TB and pulmonary TB, emphasizing the global health burden of TB. It outlines diagnostic criteria based on various cultures and histological findings, defining extra-pulmonary TB per guidelines from the American Thoracic Society and the CDC [3]. Chemotherapy protocols for pulmonary TB are reviewed, noting the importance of continued treatment to prevent drug resistance. The efficacy of rifapentine and isoniazid in the continuation phase of treatment is critically evaluated, suggesting potential benefits of once-weekly dosing

[4]. A case-control study in Samara, Russia identified key TB risk factors as raw milk consumption and unemployment. A study in Portugal linked TB risk to HIV/AIDS, prison population, unemployment, and overcrowded housing. Another study in Vitoria, Brazil highlighted the high risk of TB infection and disease among household contacts of TB patients [5]. Overall, the paper offers a comprehensive analysis of TB diagnosis, treatment, and risk factors, underlining the need for ongoing research and public health measures to combat TB globally.

III. METHODOLOGY

The stages of research methodology in applying clustering k-means and linear regression for determining the level of risk of pulmonary tuberculosis are as follows:

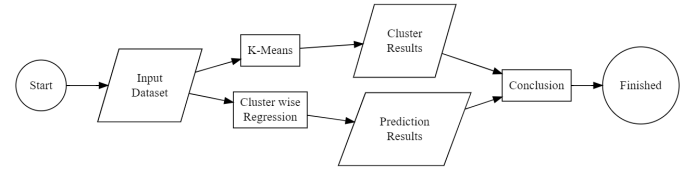


Fig. 2. Stages of Research Methodology for Determining Pulmonary TB.

The collected tuberculosis dataset is divided into two groups using k-means clustering and linear regression. The value of k will determine the number of clusters. The datasets in k-means show the cluster results. Then cluster-wise Regression is the prediction results. The combination of cluster results and prediction results is the final result. Analyzing the data in this process is more effective in determining high-risk areas for pulmonary tuberculosis.

A. Problem Identification and Data Processing

Data analysis in machine learning involves two stages: clustering model with the K-Means algorithm and Cluster Regression method to determine clusters of high-risk pulmonary TB areas and forecasting models to examine the impact of population density on the number of pulmonary tuberculosis patients and to find solutions to the problems generated based on the results of the data sets entered in the analysis.

B. Research Data Analysis

Data analysis in classifying high-risk areas for pulmonary TB. The criterion data are variables used to determine high-risk areas for pulmonary tuberculosis. The research data are as follows:

No	District	No	District	No	District
1	Dhaka	7	Gazipur	13	Mymensingh
2	Barisal	8	Jamalpur	14	Jessore
3	Khulna	9	Faridpur	15	Bogura
4	Rangpur	10	Chittagong	16	Tangail
5	Comilla	11	Sylhet	17	Dinajpur
6	Narayanganj	12	Rajshahi		

TABLE I
RESEARCH DATA

1) *K-Means Clustering Algorithm*: Data clustering is performed for each area by processing it into a cluster, and the k-means algorithm can determine cluster levels in each area based on the values of objects with different values in each group. The cluster model can also identify complex clustering values in determining high-risk areas for diseases. Furthermore, the k-means model can be divided into one or more clusters/groups. This method divides data into clusters or groups, grouping data with similar characteristics into the same cluster, and grouping data with different characteristics into other groups. The analysis of the Clustering model with the K-Means algorithm in determining risk levels in clustering can divide data into clusters into several groups. Data mining models can group data with similar characteristics into the same cluster and group data with different characteristics into another group.

- 1) Determine the number of clusters to be formed.
- 2) Decide on random centroids and initialize clusters according to the number of clusters.
- 3) Calculate the distance to the centroid using the Euclidean Distance formula, as follows:

$$d = \sqrt{(x_2 - x_1)^2 + (y_2 - y_1)^2}$$

This formula can be generalized to higher dimensions as well. In three-dimensional space, for example, with points (x_1, y_1, z_1) and (x_2, y_2, z_2) , the Euclidean distance d is:

$$d = \sqrt{(x_2 - x_1)^2 + (y_2 - y_1)^2 + (z_2 - z_1)^2}$$

And in n dimensions:

$$d = \sqrt{\sum_{i=1}^n (x_{2i} - x_{1i})^2}$$

where (x_{1i}, x_{2i}) are the coordinates of the points in each dimension.

- 4) Observe the clustering data with the closest distance value to the centroid.
- 5) Determining the data center or new centroid.

Next, Data analysis of the Linear regression model used in notation X and one response variable that can be represented by Y . Linear regression is used to assess the extent of influence between one variable and another.

2) *Cluster-wise Regression*: There are three methods for cluster-wise regression: linear regression, Finite Mixture Method (FMM), and cluster-weighted method (CWM). The linear regression method consists of one or more independent variables commonly denoted as X and one response variable represented by Y . Linear regression is used to obtain the forecast value with a variable related to another variable and can make better predictions. Therefore, it is preferable to implement a cluster initialization process using Cluster-wise Regression modeling in the next stage. One effective technique for cluster initialization is the Clustering method.

C. Research Data

The dataset for the study on the Application of K-Means Clustering and Linear Regression in Determining the Risk Level of Pulmonary Tuberculosis is as follows:

Number	Area	Population (People)	Area (km ²)	Pulmonary TB Cases
1	Dhaka	23,936,000	369	1
2	Barisal	549,000	13.23	1
3	Khulna	965,483	59.57	0
4	Rangpur	445,677	2308	6
5	Comilla	670,775	153	3
6	Narayanganj	286330	33.57	5
7	Gazipur	213 061	49.32	3
8	Jamalpur	150 172	2031.98	11
9	Faridpur	122 425	66.24	19
10	Chittagong	5,513,609	5,282.98	13
11	Sylhet	999,374	26.5	0
12	Rajshahi	983,707	34,513	18
13	Mymensingh	497,562	91.32	2
14	Jessore	110 541	2610	0
15	Bogra	944,877	72.5	5
16	Tangail	180,144	29.04	7
17	Dinajpur	206,234	20.7	1

TABLE II
RESEARCH DATA SET

IV. RESULTS AND DISCUSSION

Research utilizing the K-Means algorithm for determining the risk levels of pulmonary tuberculosis (TB) has effectively divided data into distinct clusters. This analysis categorizes areas into different risk groups: two areas fall into the first cluster (low risk), several areas into the second cluster (moderate risk), and the remaining areas into the third cluster (high risk). Following the clustering analysis, the study employs the Clusterwise Regression method to predict the impact of population density on the number of pulmonary TB cases. This approach assesses how population density influences the incidence of pulmonary TB across different areas, providing insights into the correlation between these variables and aiding in targeted public health interventions.

Application of the K-Means Algorithm Method

A. Data on the population distribution of each sub district

The population distribution in the application of k-means clustering and linear regression in determining the risk level of pulmonary tuberculosis is as follows:

No	District	Population (People)	Area (Km ²)	Number of TB Patients
1	Dhaka	23,936,000	369	1
2	Barisal	549,000	13.23	1
3	Khulna	965,483	59.57	0
...
15	Bogura	944,877	72.5	5
16	Tangail	180,144	29.04	7
17	Dinajpur	206,234	20.7	1

TABLE III
DATA ON THE DISTRIBUTION OF THE NUMBER OF PULMONARY TB PATIENTS IN EACH DISTRICT

B. Initialize the Cluster Center

The following table shows the population, area, and number of TB patients in each district, along with their assigned clusters after iteration:

Iteration	Information	Population	Area (km ²)	Number of Pulmonary TB
Dhaka	Cluster1= C1	23,936,000	369	1
Barisal	Cluster 2 = C2	549,000	13.23	1
Khulna	Cluster 3 = C3	965,483	59.57	0

TABLE IV
ITERATION DETAILS

C. Nearest Cost Values Using Euclidean Distance

Euclidean Distance is a method used to determine the shortest path or minimum distance between two points in a multidimensional space. In the context of clustering and the given table, the Euclidean Distance helps determine the nearest cluster for each data point based on their calculated cost. The nearest cost values are determined by calculating the Euclidean Distance. The results are as follows:

C1	C2	C3	Cluster Proximity	Cluster Assignment
0	23908700	23867051	0.0000	C1
23908700	0	416483.83	0.0000	C2
23867051.26	416483.0026	0	0.0000	C3
5952.07231	2645.354067	17979.00106	2645.3541	C2
4164.530169	868.3768975	19771.00014	868.3769	C2
...
24150.14635	27457.17456	48081.03475	24150.1464	C1
16433.14822	13126.26785	7498.120815	7498.1208	C3
8848.596239	5542.23089	15083.18804	5542.2309	C2
Total Proximity			98672.3064	

TABLE V
TABLE 5. DISTANCE (COST) VALUES IN THE FIRST ITERATION

D. Information

• Distance of Dhaka to Clusters

$$D_1(c_1) = \sqrt{(239636000 - 239636000)^2 + (369 - 369)^2 + (1 - 1)^2} = 0$$

$$D_1(c_2) = \sqrt{(239636000 - 549000)^2 + (369 - 13.23)^2 + (1 - 1)^2} = 239087000$$

$$D_1(c_3) = \sqrt{(239636000 - 965483)^2 + (369 - 59.57)^2 + (1 - 0)^2} = 238670517$$

• Distance of Barisal to Clusters

$$D_2(c_1) = \sqrt{(549000 - 239636000)^2 + (13.23 - 369)^2 + (1 - 1)^2} = 239087000$$

$$D_2(c_2) = \sqrt{(549000 - 549000)^2 + (13.23 - 13.23)^2 + (1 - 1)^2} = 0$$

$$D_2(c_3) = \sqrt{(549000 - 965483)^2 + (13.23 - 59.57)^2 + (1 - 0)^2} = 416483.0026$$

• Distance of Khulna to Clusters

$$D_3(c_1) = \sqrt{(965483 - 239636000)^2 + (59.57 - 369)^2 + (0 - 1)^2} = 238670517$$

$$D_3(c_2) = \sqrt{(965483 - 549000)^2 + (59.57 - 13.23)^2 + (0 - 1)^2} = 416483.83$$

$$D_3(c_3) = \sqrt{(965483 - 965483)^2 + (59.57 - 59.57)^2 + (0 - 1)^2} = 0$$

Each row in the table shows the distance of a point from each cluster center (C1, C2, C3). The "Cluster Proximity" column shows the smallest distance, and the "Nearest Cluster" column indicates the corresponding cluster assignment based on this minimum distance.

- Selection of the Fourth Centroid
- Results of the Fourth Iteration Process

The cluster centers (centroids) remain unchanged in this iteration. The new centroids obtained from the previous iteration are used to calculate the distances using Euclidean distance. The results are as follows:

Cluster	Population (people)	Area (km ²)	Number of TB Patients
Cluster 1 (C1)	55617.5	37.995	18.5
Cluster 2 (C2)	19646.33333	105.1	3.444444444
Cluster 3 (C3)	30597.16667	129.3933333	5.333333333

TABLE VI
FINAL CENTROID CENTERS FOR THE FOURTH ITERATION

These centroids represent the centers of the clusters after the fourth iteration, calculated based on the population, area, and number of TB patients. The Euclidean distance is used to determine the proximity of each district to these centroids.

E. Results of the Fourth Iteration

The calculations for the fourth iteration of applying machine learning clustering using k-means and linear regression for determining the risk level are as follows:

The results in Table 7 represent the completion of the iteration process for the third centroid. Then, the fourth iteration was carried out using the new centroids.

No	Population	Area (km ²)	No. of TB Patients	Cluster 1	Cluster 2	Cluster 3	Cluster Group
12	983707	34513	18	1704.63	37675.69	42895.36	C1
8	150172	203198	11	16410.66	19560.67	24780.32	C1
9	122425	6624	19	1704.63	34266.78	23316.17	C2
3	965483	5957	0	46376.57	10405.34	5185.74	C2
5	670775	153	3	43113.60	7142.37	1923.08	C2
13	497562	91.32	2	39703.54	3732.35	1487.30	C2
14	110541	2610	0	42924.57	6953.34	1733.85	C2
..
1	23,936,000	369	1	22445.75	13525.72	18745.46	C3
2	549,000	13.23	1	25752.77	10218.79	15438.41	C3
4	445,677	2308	6	28397.60	7573.67	12793.34	C3
6	286330	33.57	5	26607.92	9366.97	14586.03	C3
16	180,144	29.04	7	30465.50	5505.97	10725.79	C3
17	206,234	20.7	1	31293.50	4678.14	9897.47	C3

TABLE VII
RESULTS OF THE FOURTH ITERATION

The result of this process shows that there are differences in the cluster numbers for each area based on the new centroid values. In the fourth iteration, the cluster results did not change, meaning that the third and fourth clusters remained the same, and the iteration process was stopped. Therefore, it can be concluded that:

- 2 regions are included in Cluster 1
- 6 regions are included in Cluster 2
- 9 regions are included in Cluster 3

F. Results of the K-Means Clustering Algorithm

1) *Cluster Results with K-Means Using Python:* The cluster results for each sub-district, distributed across each region using the k-means algorithm, are as follows:

Sub-District	Cluster_km	Sub-District	Cluster_km
Dhaka	2	Narayanganj	2
Barisal	2	Gazipur	2
Khulna	1	Jamalpur	2
Rangpur	2	Faridpur	0
Comilla	1	Chittagong	1
Sylhet	2	Rajshahi	0
Mymensingh	1	Jessore	1
Bogra	1	Tangail	2
Dinajpur	2		

TABLE VIII
K-MEANS CLUSTER RESULTS

2) *K-Means Cluster Graph Results:* The graphical representation of the implementation of K-Means Clustering and Linear Regression in determining the risk level of pulmonary tuberculosis is as follows:

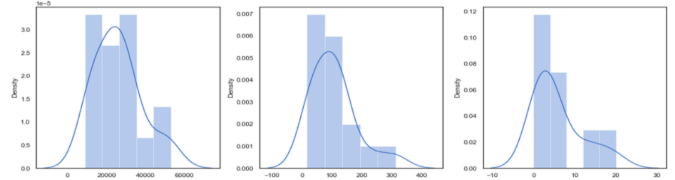


Fig. 3. K-Means Cluster Graph.

G. Linear Regression Algorithm

1) *Results of X^2 , Y^2 and XY :* The results for the values of X^2 , Y^2 , and XY to find the total are as follows:

No	X	Y	X^2	Y^2	XY
1	235	1	55225	1	235
2	192	1	36864	1	192
3	81	0	6561	0	0
4	242	6	58564	36	1452
5	98	3	9604	9	294
6	93	5	8649	25	465
7	151	3	22801	9	453
8	359	11	128881	121	3949
..
..
15	233	5	54289	25	1165
16	536	7	287296	49	3752
17	629	11	395641	121	6919
Total	9251	100	14182549	1160	109865

TABLE IX
TABLE 9. RESULTS FOR X^2 , Y^2 , AND XY

This table shows the calculated values for X^2 , Y^2 , and XY , which are used in the linear regression analysis to determine the relationship between the variables X and Y . The totals at the bottom are the sums of each column.

2) *Calculation of the Model Coefficients a and b :* To calculate the coefficients a (the intercept) and b (the slope) in the linear regression model $Y = a + bX$, we use the following formulas:

Slope (b):

$$b = \frac{n(\sum XY) - (\sum X)(\sum Y)}{n(\sum X^2) - (\sum X)^2} = 2.58415$$

Intercept (a)

$$a = \frac{(\sum Y)(\sum X^2) - (\sum X)(\sum XY)}{n(\sum X^2) - (\sum X)^2} = 0.00606$$

$$r^2 = \frac{b(\sum XY)}{\sum Y^2} = 0.57403$$

$$Y = 2.58415 + 0.00606X$$

Given the totals from Table 9:

- $\sum X = 9251$
- $\sum Y = 100$
- $\sum X^2 = 14182549$
- $\sum Y^2 = 1160$
- $\sum XY = 109865$
- $n = 17$ (number of observations)

This means that approximately 57% of the variation in the dependent variable (x), which is population density, can

explain the variation in the number of pulmonary tuberculosis patients. In other words, the variable (x) has an influence of 57% on the variable (y).

H. Calculating the Linear Regression Model Equation

The simple linear regression equation is given by:
 $Y = a + bX$. Using the calculated coefficients $a = 2.584154827$ and $b = 0.006060898$, we can predict the values of Y for given values of X .

1) *Calculation of Predicted Values:* For $X = 235$:

$$Y = 2.584154827 + 0.006060898 \times 235 = 2.584154827 + 1.424311044 = 4.008465871$$

For $X = 192$:

$$Y = 2.584154827 + 0.006060898 \times 192 = 2.584154827 + 1.163692428 = 3.747847255$$

For $X = 242$:

$$Y = 2.584154827 + 0.006060898 \times 242 = 2.584154827 + 1.46673733 = 4.050892157$$

For $X = 98$:

$$Y = 2.584154827 + 0.006060898 \times 98 = 2.584154827 + 0.594268004 = 3.178422831$$

For $X = 93$:

$$Y = 2.584154827 + 0.006060898 \times 93 = 2.584154827 + 0.563664614 = 3.147819441$$

For $X = 151$:

$$Y = 2.584154827 + 0.006060898 \times 151 = 2.584154827 + 0.915195898 = 3.499350725$$

For $X = 359$:

$$Y = 2.584154827 + 0.006060898 \times 359 = 2.584154827 + 2.176055794 = 4.760210621$$

For $X = 1138$:

$$Y = 2.584154827 + 0.006060898 \times 1138 = 2.584154827 + 6.89730199 = 9.481456817$$

By applying the regression formula, we can predict the number of tuberculosis patients based on the given population densities.

I. Graph of the Influence of Population Density on the Number of Pulmonary TB Patients

The graph depicting the influence of population density on the number of pulmonary tuberculosis patients is as follows:

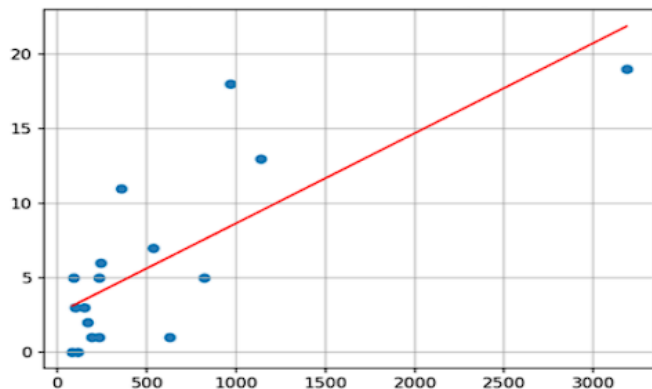


Fig. 4. Influence of Population Density on the Number of Pulmonary TB Patients.

V. CONCLUSION

The analysis using the K-means algorithm identified specific areas susceptible to pulmonary tuberculosis, guiding targeted interventions. Clustering revealed three clusters, enabling tailored strategies for each. Cluster-wise Regression showed population density explains 57% of TB variation, highlighting demographic factors' importance. However, other variables contribute to the remaining variation, warranting further exploration for comprehensive interventions. This combined approach offers a powerful framework for understanding disease patterns and informing interventions. Yet, limitations like data quality and narrow variables exist. Future research should integrate diverse data sources and employ non-linear modeling for enhanced understanding and more effective tuberculosis control strategies.

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