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Article

Enhancing Dengue Outbreak Predictions Using Machine Learning: A Comparative Analysis of Models

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Abstract: Dengue, a perilous fever-type illness transmitted by mosquitoes, remains a significant global health concern. The incidence of dengue outbreaks is primarily influenced by climate factors, contributing to fluctuating dengue cases. Therefore, this study aims to develop predictive models for dengue outbreaks employing Machine Learning (ML) techniques. Four distinct ML models, namely K-Nearest Neighbor (KNN), Random Forest (RF), Gradient Boosting Regressor (GBR), and Support Vector Regressor (SVR), were employed in this study. Prior to model analysis, thorough data preprocessing was conducted. In some instances, certain attributes exhibiting sparse correlations with dengue occurrence were eliminated based on correlation analysis. Upon finalizing the models, performance evaluation was executed through comparison based on Mean Absolute Error (MAE) metrics. The findings of this comparative analysis revealed that the KNN model exhibited significantly superior performance compared to the other three models. This outcome underscores the potential of KNN in enhancing the accuracy of dengue outbreak predictions.

Keywords: dengue; gradient boosting regressor; K-nearest neighbor; Machine Learning; Random Forest; Support Vector Regressor

I. Introduction

According to the Centre for Disease Control and Prevention (CDCP), each year, up to 400 million people get infected with dengue and approximately 100 million people get sick from infection, and 40,000 die from severe dengue. One-half of the world population is predicted to be affected in 2080 by dengue [1–5]. The virus that causes dengue fever is transmitted by the *Aedes aegypti* mosquito. Because dengue fever is spread by mosquitos, it is linked to weather and environmental factors such as temperature, precipitation, and vegetation. Increasing temperatures may worsen the problem by allowing dengue fever to spread and spread faster in low-risk or dengue-free areas of Asia, Europe, North America, and Australia. To keep patients okay, supportive care is the main thing. As a consequence, forecasting dengue epidemics is critical. With this forecast, health authorities throughout the world may take preventative actions to treat dengue fever before it spreads, potentially saving millions of lives.

Therefore, this study aims to find the most efficient model for predicting dengue cases. So—that it can prevent a dengue outbreak by accurately forecasting an increase in dengue cases. To achieve our goal—we have utilized preprocessing techniques such as the mean method to fill the null data. Correlation also has been applied to select proper features which are more connected with the dengue incidence. Then, we have used Machine Learning (ML) [6–12] methods to predict dengue cases. One more thing we have added is we worked with weather data so the predictions depend on the weather.

The rest of the study is organized as follows—Section II is all about related works which have been on the same topics. Section III contains dataset overview. Section IV contains the methodology, while Section V has the results and analysis, followed by the conclusion and future work in Section VI.

II. Related Works

The severity of the dengue outbreak has prompted various methodologies to forecast its occurrence such as [13] demonstrated the correlation between land-use factors, including agricultural land, water bodies, and forests, and reported dengue cases in Selangor, Malaysia. Employing boosted regression to accommodate non-linearities and interactions—[14] highlighted the significance of factors beyond human settlements. Subsequently, [15]—discussed the influence of human migration on dengue pathogen transmission to immunologically naive regions. Furthermore, [16] introduced a model in 2017 via Random Forest (RF) to predict dengue, diabetes, and swine flu, aiming to forecast diseases based on patient symptoms and recommend appropriate medical specialists. [17] introduced a dengue potential predicting model utilizing a dataset containing weekly dengue case records from various countries over multiple years. This model integrates meteorological parameters such as temperature, precipitation, and humidity to discern patterns and dependencies, employing Gradient Boosting Regression (GBR) to predict dengue case counts for specific weeks and years. Similarly, [18] employed RF regression—incorporating demographic, entomological, and environmental data from Singapore. Their approach achieved promising results, with over 80% of observed risk rankings falling within the predicted range. [19] explored the application of Artificial Neural Networks (ANNs) coupled with an entropy approach in Thailand to construct a prediction model for dengue epidemics. Additionally, [20] conducted in Sri Lanka and introduced an ANN model that utilized historical weather patterns and prior dengue cases to forecast epidemics in the Kandy area. These endeavors signify the diverse strategies being pursued to enhance dengue risk prediction, leveraging ML techniques and incorporating various data sources.

III. Data Analysis

In this study, we utilized data sourced from the driven data [21] repository, comprising dengue case and meteorological datasets. The dataset encompasses 1456 records, each characterized by 24 features. Key features include location, temperature, precipitation, humidity, vegetation index, alongside the total dengue cases. Notably, the dataset covers two distinct cities of Peru—Sanjuan and Iquitos, from which meteorological data is sourced, indicating the correlation between mosquito activity and dengue occurrence. This dataset, structured on a weekly basis, serves as a foundation for forecasting dengue outbreaks as shown in Table I. Sanjuan's data, spanning from 1999 to 2008, encompasses 936 records, detailing the impact of dengue cases, with Sanjuan being one of the most significantly affected cities. Conversely, the Iquitos dataset spans from 2000 to 2010, consisting of 520 recordings—reflecting a comparatively lesser impact of dengue. Our analysis focuses on both cities individually, facilitating the construction of our predictive model.

Table I. DENGUE CASES.

Data Timelines	Total Records	Feature
1999-2008 (Sun Juan)	936	24
2000-2010 (Iquitos)	520	24

IV. Methodology

In the realm of data analysis, data mining serves as the fundamental process of scrutinizing vast repositories of past data to extrapolate insights and forecast outcomes for new instances. Our study employs regression analysis as a pivotal tool to prognosticate the incidence of dengue cases in specific scenarios, leveraging a pre-existing dataset as shown in Figure 1. Initially, we conducted regression analysis comprehensively across the entire dataset. Subsequently, upon discerning correlations, we selectively pruned features exhibiting lesser correlation with the target variable. Thereafter, we iterated the regression analysis to derive enhanced predictive capabilities, facilitating an in-depth exploration of performance metrics and conducting a meticulous comparative analysis. Therefore, in the initial stage, the entire dataset undergoes a preprocessing phase aimed at ensuring data integrity.

This involves scrutinizing for any missing entries, which, if detected, are imputed with the mean value corresponding to the respective feature. Subsequently, temperature measurements, initially recorded in Kelvin, are uniformly converted to the Centigrade scale. Precision is maintained by rounding all values to three decimal places. Any essential data points are amalgamated, taking their average when necessary. Following this preparation, the dengue-related features are consolidated with the total reported dengue cases. Since the incidence of dengue in San Juan and Iquitos exhibits independence, separate dataframes are created for each locality. These dataframes are further partitioned into distinct sets for training and testing purposes. Subsequently, four distinct regression models—namely, K Neighbors Regressor (KNN), RF, GBR, and Support Vector Regressor (SVR)—are applied to the dataset for forecasting. Finally, the performance of these models is calculated via Equation (1).

$$MAE = \frac{\sum_{i=1}^n |y_i - x_i|}{n} \quad (1)$$

where n represents the total number of samples, y_i denotes the actual values, and x_i represents the predicted values. Subsequently, in the second scenario, we initiate the process by evaluating the correlations of each feature concerning the total instances of dengue cases. Following this, features exhibiting notably low or negative correlations are subjected to optimization. Post feature selection, the procedure mirrors that of the first case. Subsequent to this methodology, we conduct a comparative analysis of the outcomes to discern the optimal result and model for the specified objective.

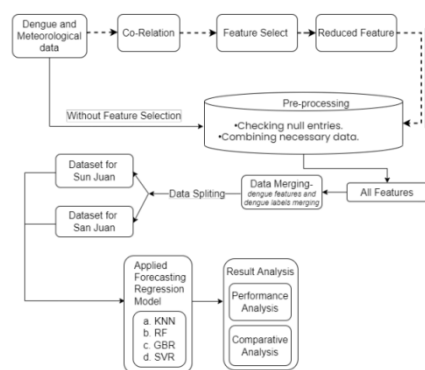


Figure 1. Flow diagram using the feature selection method.

A. Feature Selection

Understanding the relationships between different attributes within a dataset is crucial. In our analysis, the focal point is the attribute 'total case', which signifies the total instances of dengue within a specific timeframe. Our task involves assessing the correlation of each attribute with 'total case'. This correlation analysis was conducted using statistical functions available in Excel, which calculate the correlation coefficient between pairs of variables. The formula utilized for this purpose is expressed in Equation (2).

$$Correl(X, Y) = \frac{\sum(x - \bar{x})(y - \bar{y})}{\sqrt{\sum(x - \bar{x})^2 \sum(y - \bar{y})^2}} \quad (2)$$

Following the application of Equation (2), we identified certain attributes that demonstrated weaker connections with dengue occurrences than desired. Consequently, attributes exhibiting negative correlations or minimal positive correlations were excluded from further consideration. As a result, for the San Juan dataset, out of the initial 16 features, 11 were retained for experimentation as shown in Figure 2. Similarly, for the Iquitos dataset, 8 out of the 16 features were retained for experimentation as shown in Figure 3.

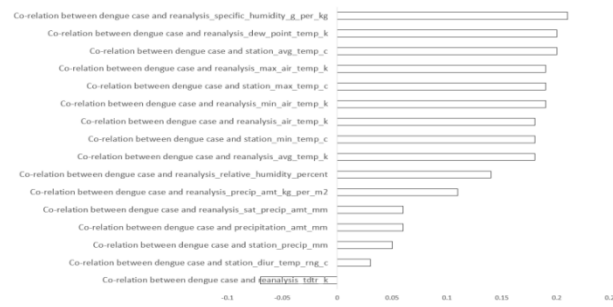


Figure 2. Correlation between all the features vs 'total case' in San Juan.

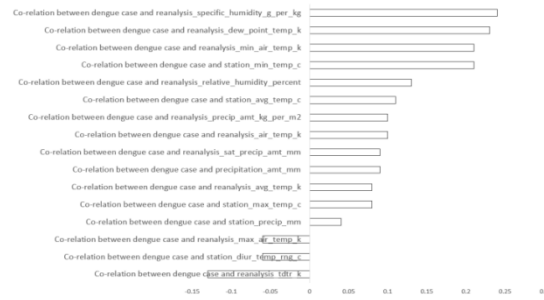


Figure 3. Correlation between all the features vs 'total case' in Iquitos.

B. Model Analysis

In our study, we undertook two preprocessing approaches for our dataset. Initially, we processed the dataset without considering correlation among its features. Subsequently, we processed the dataset accounting for correlated features. It's noteworthy that our dataset contained some missing values, the elimination of which could compromise its authenticity. Addressing missing data poses several challenges, and various techniques exist for this purpose. For our study, we opted to handle the missing data by employing a mean-based method. This method involves filling the missing values with the mean of the respective features. This approach ensures that the integrity of our dataset is maintained while mitigating the impact of missing values on our analyses. However, the concept of ML is wide and uses various algorithms which are employed to predict values or outcomes based on given data [22–25]. One such algorithm is the KNN, which determines the value of new data points by comparing them with similar points in the training set. This method relies on the assumption that data generated consistently exhibits recurring patterns. Another widely used algorithm is the RF, developed as an evolution of the bagging concept. RF combines the results of multiple decision trees to produce a final output. Despite its flexibility for tuning, RF often performs well with default settings. The GBR is another ensemble learning technique that sequentially improves upon weak decision trees. By incrementally adding models to the ensemble, GBR aims to enhance accuracy. Whereas, SVR is a supervised learning model used for classification and regression analysis. SVR aims to fit errors within a defined threshold, approximating the best value within a given margin known as the ϵ -tube. It utilizes a ϵ -insensitive loss function to penalize predictions deviating too far from the desired output [26–31].

V. Result Analysis

The experimentation in this study involved the utilization of four distinct ML model techniques as shown in Table II. Initially, employing the KNN regressor model on a preprocessed dataset yielded a MAE of 19.90 for Sanjuan and 5.52 for Iquitos. Subsequently, employing the RF model resulted in MAE values of 20.56 for Sanjuan and 5.41 for Iquitos in the second trial. Following this, the GBR was utilized, producing MAE values of 21.69 for Sanjuan and 5.39 for Iquitos. Lastly, the SVR was implemented, yielding MAE values of 26.70 for Sanjuan and 4.03 for Iquitos. Prior to feature selection, it was observed that KNN exhibited superior performance for Sanjuan, whereas SVR demonstrated better results for

Iquitos. Subsequent to feature selection, notable improvements were witnessed across both cities. KNN emerged as the optimal model for Sanjuan, achieving an MAE of 16.88, while SVR proved to be the superior choice for Iquitos, with an MAE of 4.54. For Sanjuan, MAE values of 1.34, 17.14, and 19.92 were obtained utilizing RF, GBR, and SVR models, respectively. Conversely, for Iquitos, MAE values of 5.14, 5.21, and 5.58 were recorded for KNN, RF, and GBR models, respectively. Ultimately, based on the comprehensive results obtained for both cities, KNN emerged as the most effective model for this study.

Table II. THE COMPARISON OF MAE BEFORE AND AFTER FEATURE SELECTION.

Model	Before Feature Selection		After Feature Selection	
	San Juan	Iquitos	San Juan	Iquitos
KNN	19.90	5.52	16.88	5.14
RF	20.56	5.41	17.34	5.21
GBR	21.69	5.39	17.14	5.58
SVR	26.70	4.03	19.92	4.54

A. Result Comparison

Graphical representations of Sanjuan city for each of the four models, namely KNN, SVR, and two others, are depicted in Figures 4, 5, 6, and 7, displaying both the actual and predicted values. Notably, Figure 4 illustrates a closer alignment between the actual and predicted values compared to the other figures, suggesting the superior performance of the KNN model in this context. Transitioning to the assessment of Iquitos city, Figures 8, 9, 10, and 11 offer graphical insights for each of the four models alongside the respective actual and expected values. It is evident from Figure 11 that the values exhibit greater proximity, indicating the enhanced efficacy of the SVR model for this city. Our findings are contextualized within the broader landscape of existing research, with comparisons made to prior studies utilizing the same dataset [14] as shown in Table III. Through this comparative analysis, we affirm the merit of our approach, yielding outcomes that not only meet but exceed previous benchmarks. Furthermore, our endeavor extends beyond replication, as we introduce and successfully implement a novel model, thereby achieving significant advancements in predictive accuracy.

Table III. COMPARISON BETWEEN OUR VS [14].

Model	OUR PAPER		[14]	
	San Juan	Iquitos	San Juan	Iquitos
KNN	16.88	5.14	—	—
RF	17.34	5.21	26.66	6.70
GBR	17.14	5.58	24.11	25.98
SVR	19.92	4.54	—	—

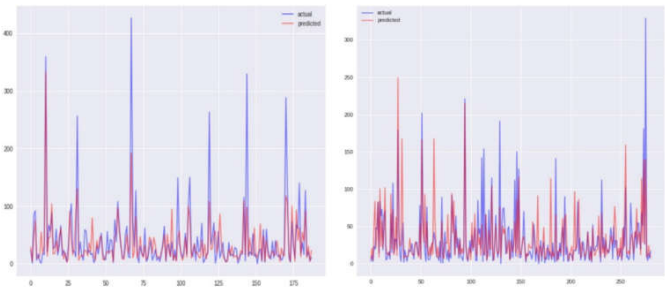


Figure 4. Comparison between true values and predicted values using KNN model before and after correlation (Sun Juan).

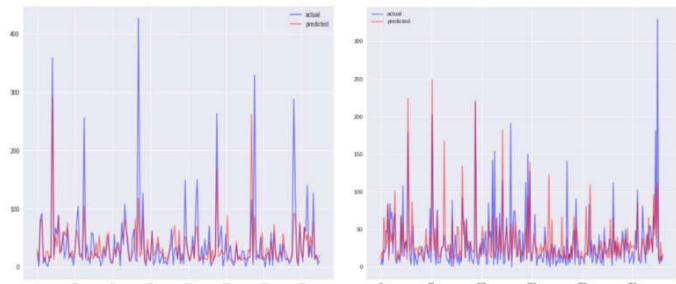


Figure 5. Comparison between true values and predicted values using RF model before and after correlation (Sun Juan).

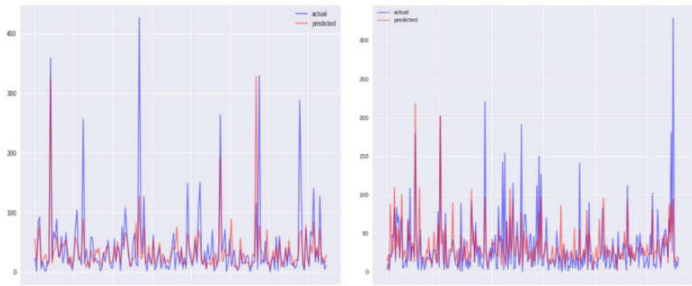


Figure 6. Comparison between true values and predicted values using GBR model before and after correlation (Sun Juan).

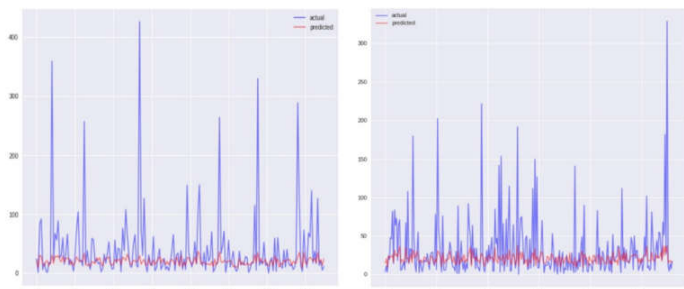


Figure 7. Comparison between true values and predicted values using SVR model before and after correlation (Sun Juan).

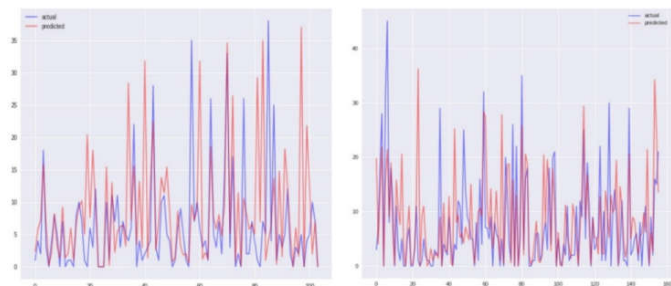


Figure 8. Comparison between true values and predicted values using KNN model before and after correlation (Iquitos).

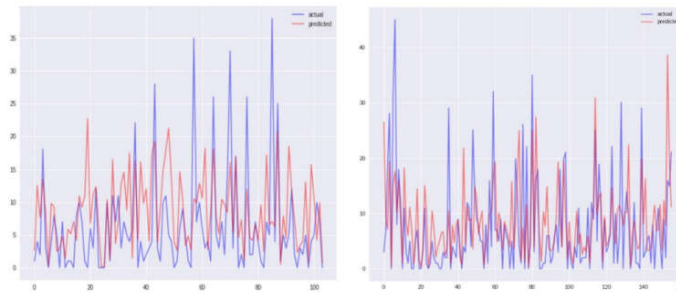


Figure 9. Comparison between true values and predicted values using RF model before and after correlation (Iquitos).

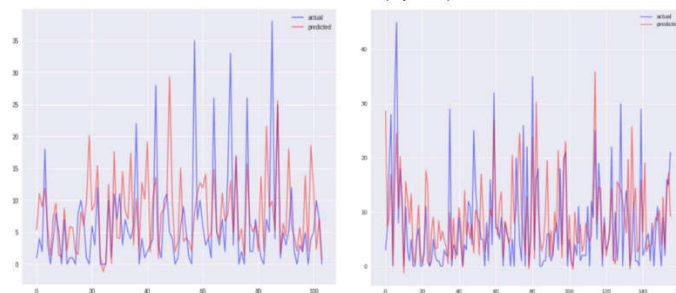


Figure 10. Comparison between true values and predicted values using GBR model before and after correlation (Iquitos).

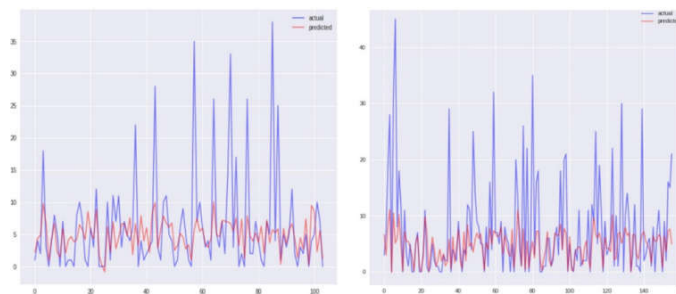


Figure 11. Comparison between true values and predicted values using SVR model before and after correlation (Iquitos).

VI. Conclusion and Future Works

In our study, we focused on analyzing a continuous dataset, which encompasses quantitative measurements including decimals and fractions, such as height, weight, and temperature. Due to the nature of our dataset, which doesn't lend itself to classification, we couldn't employ traditional classification models. Dengue, a disease affecting diverse populations, served as the focal point of our research. Leveraging ML techniques, we devised an intelligent system aimed at predicting dengue incidence based on variations in weather conditions using real-world data. Our investigation revealed a significant correlation between dengue incidence and climate variability in the selected cities. Interestingly, the primary climate factor influencing dengue incidence varied between the cities. Our findings underscore the efficacy of our forecasting model, with KNN emerging as the top performer among the evaluated models. It achieved a minimum MAE with the predominant models signifies the development of a robust forecasting framework for dengue incidence prediction.

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Data Availability: Data will be made on reasonable request.

Conflict of Interest: The authors declare that they have no known competing for financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Code Availability: Code will be made on reasonable request.

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