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The Effects of Climate on Dengue Occurrence in Matale, Sri Lanka: Generalized Linear Models

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| ARTICLE DETAILS | ABSTRACT | | |
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| Article History Received: 1 st November 2022 Accepted: 25 th September 25 Published Online: <i>31st December</i> 2024 | Dengue fever is an arboviral disease caused by four dengue virus serotypes. It is transmitted by the bite of an infected female mosquito of the genus Aedes. Over the years, dengue has become a major health issue in Sri Lanka. Since climate affects both virus incubation and mosquito development, studying the distribution of dengue fever with its relevant climate factors would be useful. This paper aims to measure the role of climate variables alone on the occurrence of dengue while eliminating the weight from past lags. The selected area for the study is | | |
| Keywords Epidemic, Dengue, Regression, Coefficient | | | |
| Email: shirinfect@gmail.com | The Matale district, and the study period is from 2005 to 2020. The engue cases were first differentiated to remove the series ependence. Monthly differentiated dengue and climate variables were og-transformed to stabilize the variance and to normalize the data sets or modeling. Cross-correlation functions (CCF) were used to measure he lag-wise correlation between dengue and climate. The causal elationship was further analyzed through a multiple-regression pproach. For that, Generalized Linear Models (GLM) were built with he Gaussian family. Two models were built considering different time ags of climate variables. The results exhibited that rainfall and average emperature positively affected dengue occurrence. Although the egression coefficients were significant, the models indicated that limate has minimal effect. However, models fitted well during epidemic periods. Adding more variables to the model to obtain more precise predictions would be useful. | | |

1. Introduction

Dengue fever (DF) and Dengue Hemorrhagic Fever (DHF) are arboviral diseases caused by four dengue virus serotypes: dengue 1, 2, 3, and 4. Aedes mosquitos, abundant in tropical and subtropical areas worldwide, transmit the virus. Dengue fever has spread substantially over the last few decades, and the disease is now endemic in various parts of the world [1].

Dengue has been endemic in Sri Lanka for the last 40 years. Since the early 1990s, significant epidemics of severe and fatal DHF have occurred at regular intervals. The magnitude of DF and DHF epidemics in Sri Lanka continued to increase during the 2000s, with a major outbreak in 2009 of 35,008 suspected cases [2]. Dengue became a serious public health issue in the country between 2010 and 2016, with cases progressively increasing. In 2017, the Central Epidemiology Unit of the Ministry of Health in Sri Lanka received reports of 186,101 suspected cases and 440 dengue-related deaths. This is the highest amount of cases reported in a year in Sri Lanka since dengue fever was declared a notifiable disease in 1996. Dengue fever became a serious public health issue in the United States between 2010 and 2016, with cases progressively increasing (from 28,473 in 2011 to 55,150 in 2016) across the country [3]. In 2019, the second-greatest cases (105,049 dengue cases) were reported. Initially, dengue fever spread in urban regions but has since expanded to rural and hilly countries.

Numerous studies have discovered the and relationship between climate dengue transmission. Increased temperature can promote dengue risk by improving vector development rate and shortening the viral incubation period. On the other hand, extreme hot temperatures reduce dengue transmission by increasing the mosquito mortality rate. Rainfall may provide a habitat for Aedes Aegypti and Aedes Albopictus larvae and pupae, but heavy rainfall will flush them away [5]. As a result, studies on the relationship between climate variables and dengue are important for deciding dengue outbreaks.

This paper aimed to determine the impacts of climatic conditions on dengue transmission in a selected district of Sri Lanka. A lag-wise correlation was observed between climate variables and dengue cases. The critical climate variables were recognized with their affecting time lags. Thereafter, the degree of their influence was assessed using a Generalized Linear Model (GLM). In addition, the association between climate and dengue epidemic was measured by fitting the model on dengue outbreaks. This study estimates the effects of climate alone, which is valuable for evaluating its function on different dengue patterns.

2. Material and Methods

2.1. Study area

The research was conducted in the Matale District of the central province of Sri Lanka. The land area of Matale is 1,993 km² and is divided into 11 divisional Secretariats divisions. The estimated population of the area was 484,531 in 2012, including urban, semiurban, and rural areas [12]. Matale is the secondhighest dengue-affected area in the central province.

2.2. Data

Monthly dengue cases from 2005 to 2020 were obtained from the Regional Director of Health Services (RDHS) in Matale with the permission of the Provincial Director of Health Services in the Central Province. Climatic data was obtained from the Department of Meteorology of Sri Lanka.

2.3. Methodology

Monthly aggregated dengue and climatic variables were used to analyze the association between dengue cases and climate. Rainfall, minimum temperature, maximum temperature, average temperature, and humidity were considered climatic variables. Dengue cases were first differentiated to remove the series' dependence on time. Then, the variables were log-transformed to stabilize the variance and to normalize the data sets for the modeling [6].

Pearson correlation coefficient was used in crosscorrelation functions, and lagged correlations between dengue and climate variables were observed. A multiple regression approach examined the impact of climatic variables on dengue cases. For that, Generalized Linear Models (GLM) were built with the Gaussian family. GLM consists of a random component specifying the conditional distribution of the response variable, Yi (for the ith lag of independently sampled dengue cases), given the values of the explanatory variables in the model. Yi and explanatory variables are the members of the Gaussian family that are continuous, decimal data with normal distribution [7,8].

The monthly lags of climate variables used as explanatory variables were determined from crosscorrelation functions (CCF). We built two models, and the best model was determined based on the lowest Akaike's information criterion (AIC), Bayesian information criterion (BIC), and residual sum of squares (RSS) [9].

Model-1: A generalized Linear Model (GLM) was fitted using log₁₀ (differentiated cases) as the response variable and log₁₀ (climatic variables) with only the highest correlated lagged terms as the explanatory variables.

Model-2: All the lagged terms of climate variables were considered up to three lags before eliminating regression coefficients through significance. The final model was determined with significant coefficients.

For a more detailed study, the mean plus two standard deviation method was used to detect abnormal dengue transmission periods by modifying the recommendations of Badurdeen et al. [13,14]. Moreover, model-1 and model-2 were fitted again on the transformed dengue observations from abnormal transmission periods and with their respective climate variables. The R-squared values are used to compare the models.

3. Results and Discussion

Among climate variables, rainfall and average temperature contributed to dengue cases in the Matale region. Significantly correlated lags were identified through cross-correlation functions for rainfall and average temperature. The rainfall has a significant positive (p<0.05 at 95% confidence interval) association in 1-2-month lags while the temperature reached in 2-3-month lags.



Figure 1: CCF- dengue cases and rainfall

Figures 1 and 2 show that lag 1 of rainfall and lag 2 of average temperature are the highest correlated lagged terms. Therefore, the GLM equation for model-1 was,

$$Y_t = \beta_0 + \beta_1 (RF)_{t-1} + \beta_2 (Temp)_{t-2}$$





Figure 2: CCF- dengue cases and temperature

Where; Y_t is the predicted value of dengue observation at the time *t* and β_j is regression coefficient. RF_{t-j} is the log₁₀ (Rainfall) at time lag *j* and $Temp_{t-j}$ is the log₁₀(Temperature) at time lag *j*.

The results of model 1 are summarized in Figure 3. Figure 4 illustrates the results of model-2. After eliminating insignificant coefficients, the final model includes rainfall lags 1, 2, and temperature lag 2. The values of AIC, BIC, and RSS are summarized in Table 1.

Call: glm(formula = Cases ~ RF_Lag1 + Temp_Lag2)

Deviance Residuals: Min 10 Median 30 Max -2.7331 -0.9325 0.1496 0.9345 2.9668 Coefficients: Estimate Std. Error t value Pr(>|t|) (Intercept) -18.7909 5.9717 -3.147 0.00192 ** RF_Lag1 0.8291 0.1792 4.626 6.93e-06 *** Temp_Lag2 12.0175 4.1129 2.922 0.00391 ** ---

Signif. codes: 0 (****' 0.001 (**' 0.01 (*' 0.05 (.' 0.1 (' 1 (Dispersion parameter for gaussian family taken to be 1.4467)

Null deviance: 307.70 on 190 degrees of freedom

Residual deviance: 271.98 on 188 degrees of freedom (2 observations deleted due to missingness) AIC: 617.54

Number of Fisher Scoring iterations: 2

Figure 3: Output of model-1

 $Y_t = 0.8291(RF)_{t-1} + 12.0175(Temp)_{t-2} - 18.7909$

is the predicted regression equation of model-1. All the regression coefficients are significant at a 95% confidence interval with p<0.05.

$$\begin{split} Y_t &= 0.7324 (RF)_{t-1} + 0.5577 (RF)_{t-2} + \\ 17.2274 (Temp)_{t-2} - 27.1462 \end{split}$$

is the predicted regression equation of model 2. Rainfall at a time lag of 1 is the most critical climate variable with the smallest p-value and standard error. Call: glm(formula = Cases ~ RF_Lag1 + RF_Lag2 + Temp_Lag2) Deviance Residuals: 3Q 0.8827 Min -2.5077 1Q -0.9368 Median 0.1958 2.8195 Coefficients: Estimate Std. Error t -27.1462 6.5068 -0.7324 0.1797 value Pr(>|t|) -4.172 4.63e-05 4.076 6.79e-05 (Intercept) RF Lag1 RF_Lag2 0.557 0.1924 2.898 0.004205 17,2274 4.4057 3.910 0.000129 Temp_Lag2 Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 (Dispersion parameter for gaussian family taken to be 1.396095) Null deviance: 307.62 on 189 degrees of freedom Residual deviance: 259.67 on 186 degrees of freedom (3 observations deleted due to missingness) AIC: 608.55

Figure 4: Output of model-2

Table 1: Measures of model performance

Number of Fisher Scoring iterations: 2

| Model | AIC | BIC | RSS |
|-------|--------|---------|----------|
| 1 | 617.54 | 630.533 | 271.9796 |
| 2 | 608.5 | 624.778 | 259.6736 |

Considering the lowest AIC, BIC, and RSS values, model 2 was selected. Figure 5 shows the distribution log₁₀ (differentiated cases) predicted by rainfall and temperature. We were able to capture the pattern of dengue cases from the predicted values. The graph indicates that the climate variables alone are insufficient to detail all the information on dengue cases. The corresponding R-squared value of the model equals 0.16. Although we obtained a lower Rsquared value, the smaller p-values for the estimates indicate the significant association between climate variables and dengue transmission.



Figure 5: Actual dengue observations with fitted values. Red line represents fitted value and black line represents actual dengue observations.

The role of climate on the dengue outbreaks was evaluated after detecting epidemic periods. The highest correlated lags were obtained at rainfall lag1 and temperature lag2 on abnormal transmissions. $Y_t = 2.2805(RF)_{t-1} + 58.5671(Temp)_{t-2} - 88.3988$

is the new predicted equation from model-1 considering extreme dengue cases alone. The respective R-squared value is 0.38. All the regression coefficients are significant here.

The refitted model-2 on abnormal transmission results in the following equation,

 $Y_t = 2.7733(RF)_{t-1} + 2.0224(RF)_{t-2} + 82.2899(Temp)_{t-2}$

The R-squared value is 0.57, which explains 57% of information on dengue observation. As previously stated, it was only 16% when the entire series of dengue observations was included.

4. Conclusion

Dengue has been a critical public health problem in Matale and other parts of Sri Lanka. This study aimed to find a statistical method to ensure the role of climate variables on the occurrence of dengue while eliminating the weight from past lags. Generalized Linear Modeling is a powerful tool to explore the causal relationship between independent dengue observations and explanatory climate variables. The results of the models show that the number of dengue cases increases as rainfall and average temperature increases. The rainfall impacts dengue fever in 1-2 months and average temperature within 2-3 months. The degree of effect was minimal when we considered the entire dengue series from the study period. The impact was higher when it came to epidemic periods. Aside from predicting future dengue cases based on climate variables, we attempted to study the influence of climate alone on various dengue patterns, such as normal versus atypical transmissions. Adding more variables to the model to obtain more precise predictions would be useful.

References

- Schaefer TJ, Panda PK, Wolford RW. Dengue Fever. [Updated 2022 Apr 22]. In: StatPearls [Internet]. Treasure Island (FL): StatPearls Publishing; 2022 Jan-. Available from: https://www.ncbi.nlm.nih.gov/books/NBK430732/ [Accessed 10th September 2022].
- Ministry of Health, Nutrition and Indigenous Medicine. National Dengue Control Unit - Sri Lanka Situation, Available from: http://www.dengue.health.gov.lk/web/index.php/en/i nformation/sri-lanka-situation [Accessed 12th September 2022].

- Tissera HA, Jayamanne BDW, Raut R, Janaki SMD, Tozan Y, Samaraweera PC, Liyanage P, Ghouse A, Rodrigo C, de Silva AM, Fernando SD. Severe Dengue Epidemic, Sri Lanka, 2017. Emerg Infect Dis. 2020 Apr;26(4):682-691. Available from: doi:10.3201/eid2604.190435. PMID: 32186490; PMCID: PMC7101108.
- Department of Health. Living well with dementia: a national dementia strategy. Available from: https://www.gov.uk/government/publications/livingwell-withdementia-strategy [Accessed 4th June 2015].
- Naish, S., Dale, P., Mackenzie, J.S. et al. Climate change and dengue: a critical and systematic review of quantitative modelling approaches. BMC Infect Dis 14, 167 (2014). Available from: https://doi.org/10.1186/1471-2334-14-167 [Accessed 10th September 2022].
- S Shrestha MB, Bhatta GR. Selecting appropriate methodological framework for time series data analysis. The Journal of Finance and Data Science. 2018 Jun 1;4(2):71-89.
- Hardin JW, Hardin JW, Hilbe JM, Hilbe J. Generalized linear models and extensions. Stata press; 2007.
- Fox J. Applied regression analysis and generalized linear models. Sage Publications; 2015 Mar 18.
- Withanage GP, Viswakula SD, Nilmini Silva Gunawardena YI, Hapugoda MD. A forecasting model for dengue incidence in the District of Gampaha, Sri Lanka. Parasites & vectors. 2018 Dec;11(1):1-0.
- Pears R, Shields G. Cite them right: the essential referencing guide. Palgrave study skills. 10th ed. Basingstoke: Palgrave; 2016.
- Ramalho R, Helffrich G, Schmidt DN, Vance D. Tracers of uplift and subsidence in the Cape Verde archipelago. Journal of the Geological Society. 2010;167(3): 519–538. c doi:10.1144/0016-76492009-056.
- Department of Census and Statistics. Census of Population and Housing - 2012. Available from: http://www.statistics.gov.lk/PopHouSat/CPH2012Vis ualization/htdocs/index.php?usecase=indicator&acti on=DSMap&indId=10&district=Matale&Legend=3 [Accessed September 30, 2022].
- Badurdeen S, Valladares DB, Farrar J, Gozzer E, Kroeger A, Kuswara N, Ranzinger SR, Tinh HT, Leite P, Mahendradhata Y, Skewes R. Sharing experiences: towards an evidence based model of dengue surveillance and outbreak response in Latin America and Asia. BMC Public Health. 2013 Dec;13(1):1-4.

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14. Udayanga L, Aryaprema S, Gunathilaka N, Iqbal MC, Fernando T, Abeyewickreme W. Larval indices of vector mosquitoes as predictors of dengue epidemics: an approach to manage dengue outbreaks based on entomological parameters in the districts of Colombo and Kandy, Sri Lanka, BioMed Research International 2020 June 16; 2020