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Spatial multi-layer perceptron model for predicting dengue fever outbreaks in surabaya(Article)([Open Access](#))

Halim, S., Handojo, A., Widodo, I.E., Felecia, Octavia, T.

^aPetra Christian University, Industrial Engineering Department, Surabaya, 60238, Indonesia^bPetra Christian University, Informatics Engineering Department, Surabaya, 60238, Indonesia

Abstract

Dengue fever (DF) is a tropical disease spread by mosquitoes of the Aedes type. Therefore, a DF outbreak needs to be predicted to minimize the spread and death caused by it. The spread of dengue fever is a spatial problem. In this paper, we adopted the Multi Linear Perceptron (MLP) to solve the spatial problem, and we called it a spatial multi-layer perceptron model (Spatial MLP). In this proposed model, we consider two types of input neurons in the Spatial MLP, a region and the neighbourhood of that region. The spatial inputs dynamically change to the region. Additionally, the neighbourhood numbers of a region are also varied. So, the spatial inputs are changed in terms of the number of inputs and the neighbourhoods. As a result, the proposed model is outperformed the traditional MLP since it can adapt to the neighbourhoods. We can conclude the spatial MLP model can manage the information and predict the dengue fever outbreak in Surabaya © 2020 ASTES Publishers. All rights reserved.

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Spatial Multi-Layer Perceptron Model for Predicting Dengue Fever Outbreaks in Surabaya

Siana Halim^{1,*}, Andreas Handojo², Ivan Enrico Widodo², Felecia¹, Tanti Octavia¹

¹Petra Christian University, Industrial Engineering Department, Surabaya 60238, Indonesia

²Petra Christian University, Informatics Engineering Department, Surabaya 60238, Indonesia

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ABSTRACT

Dengue fever (DF) is a tropical disease spread by mosquitoes of the *Aedes* type. Therefore, a DF outbreak needs to be predicted to minimize the spread and death caused by it. The spread of dengue fever is a spatial problem. In this paper, we adopted the Multi Linear Perceptron (MLP) to solve the spatial problem, and we called it a spatial multi-layer perceptron model (Spatial MLP). In this proposed model, we consider two types of input neurons in the Spatial MLP, a region and the neighbourhood of that region. The spatial inputs dynamically change to the region. Additionally, the neighbourhood numbers of a region are also varied. So, the spatial inputs are changed in terms of the number of inputs and the neighbourhoods. As a result, the proposed model is outperformed the traditional MLP since it can adapt to the neighbourhoods. We can conclude the spatial MLP model can manage the information and predict the dengue fever outbreak in Surabaya

1. Introduction

Dengue Fever (DF) outbreak happened annually, but every year the number of victims is very high. In the present decade, Ketharpal mentioned that dengue is endemic to 128 countries, mostly developing nations, posing a risk of death to approximately 3.97 billion people annually [1]. Cartographic approaches estimated that 390 million dengue infections annually, out of which 96 million cases evident apparently [2,3]. World health organization (WHO) stated that more than 70% of the population at risk for dengue worldwide live in member states of the WHO South-East Asia Region and Western Pacific Region [4]. WHO categorized the variable endemicity of dengue fever into four categories. Indonesia is included in category A which means the endemic occurs due significant public health problem, a leading cause of hospitalization and death among children, hyperendemicity with all four serotypes circulating in urban areas, and spreading to rural areas [5].

More than thirty-nine thousand (39,876) DF cases and 254 deaths were reported by Indonesian Health Ministry from January to March 2020 [6,7,8]. For significant reduction of dengue mortality, the strategies for the prevention of dengue include prompt diagnosis of fever cases, providing appropriate clinical management, and controlling vector, and personal protection methods. Therefore, severe cases can be managed with appropriate treatment, and health personnel at all level can be trained. Improved outbreak prediction and detection through coordinated surveillance will be able to reduce DF spread and effected area [9, 10].

Many types of research have been done in predicting the spread and DF affected area. A five years dataset from Sleman, a district in Central Java Indonesia are used for predicting the spread of the DF [11]. Mahdiana's model is based on vector autoregressive spatial autocorrelation (varsa). A four years dataset from Bandung stated that the incidence rate of dengue fever was not related to annual rainfall, population density, larva free index, and prevention Program [12]. The spreading of DF in Surabaya, Indonesia, is modelled using statistical learning [13,14]. Besides of statistical learning approach, many researchers also developed the model in the machine learning approach. Various machine learning algorithms are compared, such as naive Bayes, random forests, minimal sequential optimization [15]. They collected data from the health department, Karuna medical hospital, Kerala, and online sources. The authors stated random forests gives better accuracy for the early detection of dengue disease.

On the other hand, the use of neural networks as an algorithm for predicting disease has been widely used. An artificial neural network is used to predict the DF outbreak in Srilanka [16], using similar approach [17,18] the DF outbreak in Thailand modelled and in the Northwest Coast of Yucatan, Mexico and San Juan, Puerto Rico, respectively. Most of the artificial neural network that has been used to develop the model is multi-layer perceptron, with the input as the population characteristics in each region and number of DF infected in the previous years for predicting the number of DF infected in the current or next year.

*Corresponding Author: Siana Halim, Email: halim@petra.ac.id

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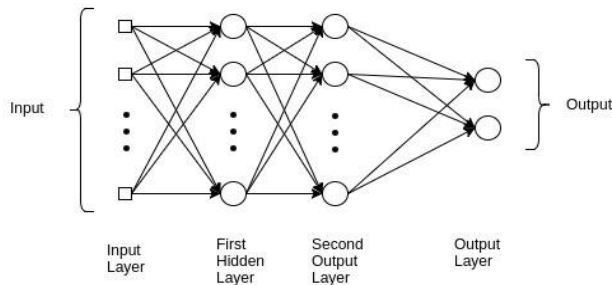


Figure 1. Multi-Layer Perceptron Neural Network

In this research, we proposed a spatial multi-layer perceptron model for predicting the DF outbreak. As a case study, we used DF data in Surabaya. The proposed model adopts the spatial approach in statistical learning as well as the multi-layer perceptron in the machine learning approach. This proposed model tries to accommodate the nature of DF disease spreading. Because DF is a type of disease that spreads through dengue mosquitoes, if DF infects a particular area, the surrounding areas will be vulnerable to the spread of the disease (spatially correlated). Therefore, disease prediction in a particular area is greatly influenced by the DF disease in the surrounding area. So, to predict the possibility of the spread of dengue fever in a particular area, we need to calculate the spread DF data from the surrounding areas. This data will be calculated separately for each region (spatial dependent). This proposed model will implement in Multi-Layer Perceptron Model (MLP) Neural Network. The MLP NN model does not accommodate the spatial dependency in the neural-network construction. This proposed model tries to build a spatial MLP model to accommodate the nature of DF disease spreading.

Additionally, we also present the model for predicting the DF web basely. Since currently, the data for DF victims is manually collected at community health centers, and it will be reported to the regional health department. Based on this DF data, the city and province will take a curative and preventive action to prevent DF outbreak in next year. Urgent measures also being taken by community health centers during outbreaks such as fogging or spreading abate powder in water collecting area. Without a sound information system on DF outbreak location and spreading, the government cannot control and minimize dengue mortality.

2. Research Methods

2.1. Multilayer Perceptron

Multilayer perceptron (MLP), also often called as feedforward neural networks consists of neurons that are ordered into layers (Figure 1). The first layer is called the input layer, and the last one is called as the output layer, the layers between are hidden layers [19].

The main goal of MLP is to approximate some function f ; e.g. in a regression, $Y = \theta_0 + \theta_1 X_1 + \dots + \theta_n X_n$; the function $Y = f(X)$ maps the input vector X into the a value Y . The feedforward network defines a mapping $Y = f(X; \theta)$ and learns the value of the parameters θ that result in the best function approximation.

In the general MLP (Figure 1), we know that each layer can be modelled as a function of

$$Y = f(\theta X + b) \quad (1)$$

where f is the activation function, θ are weights in the layer, X is the input vector, which can also be the output of the previous layer, and b is the bias vector. The hidden layers, which are located in between the input and the output of a neural network, will perform nonlinear transformations of the input in the network. The number of the hidden layers are varied. It depends of the function of the neural network. Similarly, the number of the layers may vary. It depends on their associate weights [20].

The function f is called the transfer function. The transfer function used in this research is ReLu (Rectified Linear Unit) [21]. This function is defined as $y = \max(0, X)$. Visually it can be seen in Figure 2.

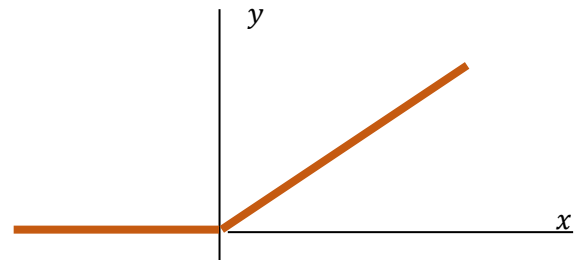


Figure 2. Rectified Linear Unit

2.2 Spatial Multilayer Perceptron

It is well known that the dengue fever happening most in tropical countries and considered as the fastest spreading mosquito-borne disease. It is transmitted by Aedes mosquito which infected with a dengue virus. The spreading of this diseases is spatially correlated [13]. The MLP model does not accommodate the spatial dependent in the neural-network construction. Therefore, in this paper we modified the Multilayer Perceptron Model (MLP), to accommodate the spatial nature of the disease.

In this model, we assumed that the spread of the diseases is in the first-order contiguity level. That is, the number of cases in location s is contagious to its north, east, south, and west neighborhoods. Some additional explanatory variables are also included in the model. They are sex ratio, poverty percentage, population density. In this proposed model, the first layer equation can be written as follows:

$$Y(s) = f\left(\sum_{j=1}^N \left(\sum_{i=1}^M w_{ij}^1 X_i(s) + \sum_{k \in (nr, es, st, wt)} w_j^2(s_k) Y(s_k)\right) + \varepsilon\right) \quad (2)$$

where:

- $Y(s)$: Number of cases in the location s
- w_{ij}^1 : The j th neuron weight w.r.t explanatory variable i
- $X_i(s)$: The explanatory variable i in the location s
- $w_j^2(s_k)$: The j -th neuron weight w.r.t response variable Y in the location s_k
- $Y(s_k)$: Number of dengue fever cases in the location s_k
- s_k : The k (north, east, south, west) location of the location s
- ε : Bias

M : Number of explanatory variables ($M = 3$)
 N : Number of neurons
 i, j, k : index

In this model the input of the MLP is changed depend on the location s . To give an illustration, let predicts the number of cases in sub-district Balongsari (Figure 3). This region shares borders to sub-district Asemrowo (north), sub-district Tanjungsari (east), sub-district Lontar (south) and sub-district Manukan Kulon (west).

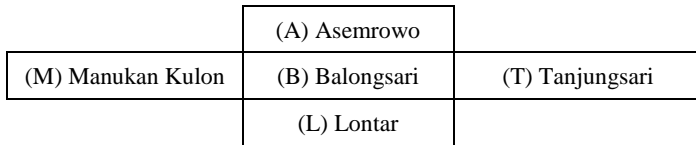


Figure 3. Sub-district Balongsari surrounding area

So, the model can be written as

$$Y(B) = f(\sum_{j=1}^N w_{1j}^1 X_1(B) + w_{2j}^1 X_2(B) + w_{3j}^1 X_3(B) + w_1^2(A)Y(A) + w_2^2(T)Y(T) + w_3^2(L)Y(L) + w_4^2(M)Y(M))$$

here $X_1(B)$ is the sex ratio in Balongsari, $X_2(B)$ is the poverty percentage in Balongsari, $X_3(B)$ is the population density in Balongsari. $Y(A), Y(T), Y(L)$ and $Y(M)$ are the number of DF case in Asemrowo, Tanjungsari, Lontar and Manukan Kulon respectively. The input neurons of the model adaptively changes with respect to the region s .

2.3 Design Spatial Multi-Layer Perceptron Neural Network

The design uses seven neurons; three neurons represent sex ratio, percentage of poverty and population density of each region under health community center s recorded in 2018. The other four neurons are dynamic neurons. They represent the number of cases in the north, east, south, and west. These neurons depend on the location s (See Figure 4).

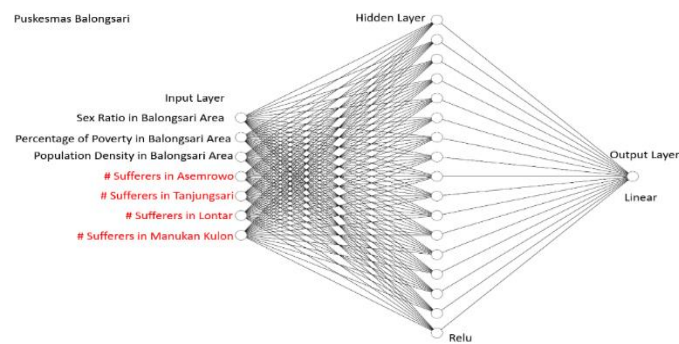


Figure 4. Illustration for sub-district Balongsari

We used 252 data training (data from 2012-2015) and 126 (data from 2017-2018) data testing. The training process used 3500 epochs (Table 1), and mean squared error is used to measure the loss/error function and we used the stochastic gradient descent as the optimizer.

After some modeling the best design for this case used 1 hidden layer with 17 neurons and 1 output layer (Table 2). The activation function is rectified linear unit (Relu) on the hidden

layer and linear on the output layer (Figure 5). This model is implemented as Python functions. It can be used to the other regions as far as the dataset is provided.

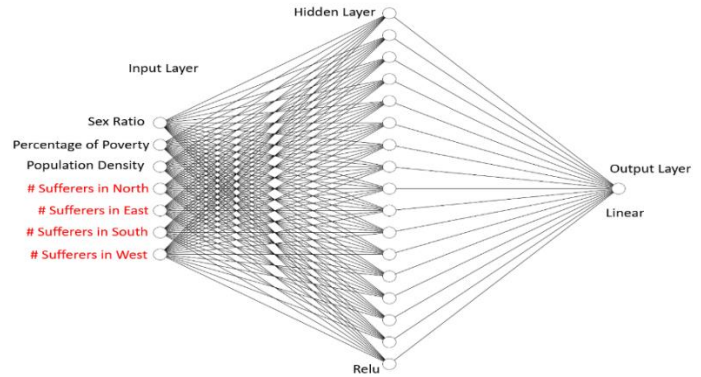


Figure 5. The design of spatial MLP model

Table 1. Setting the Number of Epochs and Neuron

Model	Epochs	Neuron	Loss on Data Training	Loss on Data Validation
1	1500	15	0.0198	0.0403
2		17	0.0187	0.0407
3		20	0.0196	0.039
4		22	0.0182	0.039
5	3500	15	0.0189	0.0304
6		17	0.0176	0.0294
7		20	0.0184	0.0383
8		22	0.0171	0.0399
9	4000	15	0.0184	0.0309
10		17	0.0171	0.0304
11		20	0.0181	0.0381
12		22	0.0169	0.0302

Table 2. Setting the Hidden Layer

Model	Layer	#Neuron Hidden Layer 1	#Neuron Hidden Layer 2	Loss on Data Training	Loss on Data Validation
1 hidden	1	17	-	0.0176	0.0294
2 hidden	2	17	7	0.0181	0.0297

3. Result and Discussion

3.1. Data Collection

Data we collected from Surabaya city consist of weather and population characteristic data. Weather data records the number of rainy days in a year, precipitation, maximum and minimum temperature, maximum and minimum humidity. The result shows that Surabaya weather is not significantly different, so it will not be used as the model's explanatory factor. Population characteristic data will be used in the model, and they are sex ratio, population density, and poverty percentage.

3.2. Data Training and Testing

We use the recorded data from 2012-2015 as the training dataset and the data from 2016 to validate the model. The training dataset consists of $63 \times 4 = 252$ data. As usual, we normalized the data set in advanced. The loss of the training data is 0.0176. Figure

5 shows the fitting of the real data to the prediction one. The horizontal axe represents the community health center, the vertical axe represents the number of cases in each community health center, recorded from 2012-2015. Figure 6 shows that the prediction can follow the pattern of the real dataset. During 2012-2013 the number of cases was high, and it started to drop in 2014-2015. The box plot of the data training (Figure 7) shows that there are several outliers in the real dataset and those outliers cannot be captured by the proposed model. The median of the prediction is not significantly different from the real one, but the interquartile range of the prediction is smaller than the real dataset. The two-samples t-test for the training data set is summarized in Table 4. The one-sided p-value is 0.335, we can conclude that there is no mean difference between the real dataset and the predicted one. The mean difference is -0.53 and the 95% confidence interval of the mean difference is (-2.96, 1.91).

Table 3. Surabaya Statistics in 2018

	Min	Mean	Max
Population (thousand)	12541	45802	87561
Area (Km2)	0.915	2.001	14.400
Density (thousand/Km2)	2733	46992	541022
Sex Ratio (Men/Women)	91.5	99.27	110.93
Poverty percentage (%)	4.03	18.02	55.46
Rainy day (days/month)	9.83	13.99	16.00
Precipitation (mm/month)	129.9	164.6	194.9
Max Humidity per month	70	88.72	94.75
Min Humidity per month	46.08	53.14	57.83
Max Temperature	28.21	33.30	34.43
Min Temperature	23.11	26.29	28.73

Table 4. Two-samples t-test for Data Training

	Real	Predicted
Mean	18.86508	19.39285714
Variance	262.0455	125.4745304
Observations	252	252
Hypothesized Mean Difference	0	
df	447	
t Stat	-0.425602637	
P(T<=t) one-tail	0.335301126	
t Critical one-tail	1.648269625	
P(T<=t) two-tail	0.670602253	
t Critical two-tail	1.965285234	



Figure 6. The real and prediction line chart of data training from 2012-2015.

We use the recorded data from 2017-2018. There are 126 data. Applying the modelled, the loss value of the testing dataset is 0.052. Some of the prediction are lower/higher than the reality (see Figure 8). Some community health centers reported that there

were no dengue fever cases in their area (the number of cases equal to zero), but in their surrounded areas reported highly dengue fever cases. As the result, the real zero number cannot be captured as zero in the model. The model will predict the number of infected in that area as the mean value of the neighborhood. This situation is acceptable, since the predicted number will give early warning to that region to prevent the outbreak in that area.

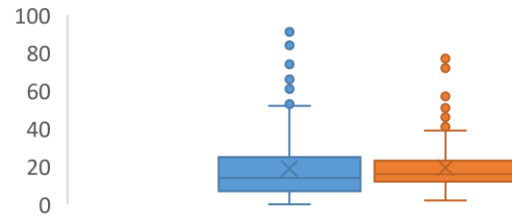


Figure 7. Box plot of the training dataset

3.3 Discussion

In this study, we proposed a spatial-MLP model, which accommodate the spatial property of the dataset. Comparing the other NN models which are used by [16,17,18,19,20] this model uses dynamic variables, which depend on the neighborhood of a region as well as the external variables. The model by [16,17,18,19,20] used only external variables, which do not depend on the neighborhood of a region.

The same dataset has been modelled using the Geostatistical Weighted Regression (GWR) [13]. In [13], the predicted model can follow the pattern of the actual dataset. However, the MSE of the prediction for the years 2017-2018 is 8.59. Compare to this model, the mean square error of the testing data set in the same years is lower, that is 8.07. The t-test shows that there is no significant difference between the mean of the real and the mean of predicted of the testing dataset (Table 5). This model is better than the GWR. However, the computation time of GWR is faster than the spatial-MLP. Since in the spatial-MLP, we have to do the hyperparameter tuning for finding the best model. This model has limitation. It cannot capture the zero in the dataset. The zeros will be predicted as the mean values of the surrounding areas. In this study we proposed a spatial-MLP model, which accommodate the spatial property of the dataset. Comparing the other NN models which are used by [16,17,18] this model uses dynamic variables, which depend on the neighborhood of a region as well as the external variables. The model by [16,17,18] used only external variables, which are not depend on the neighborhood of a region.

The same dataset has been modelled using the Geostatistical Weighted Regression (GWR) [13]. In [13], the predicted model can follow the pattern of the true dataset. However, the MSE of the prediction for the years 2017-2018 is 8.59. Compare to this model, the mean square error of the testing data set in the same years is lower, that is 8.07. The t-test shows that there is no significant difference between the mean of the real and the mean of predicted of the testing dataset (Table 5). This model is better than the GWR. However, the computation time of GWR is faster

than the spatial-MLP. Since in the spatial-MLP, we have to do the hyperparameter tuning for finding the best model. This model has limitation. It cannot capture the zero in the dataset. The zeros will be predicted as the mean values of the surrounding areas.

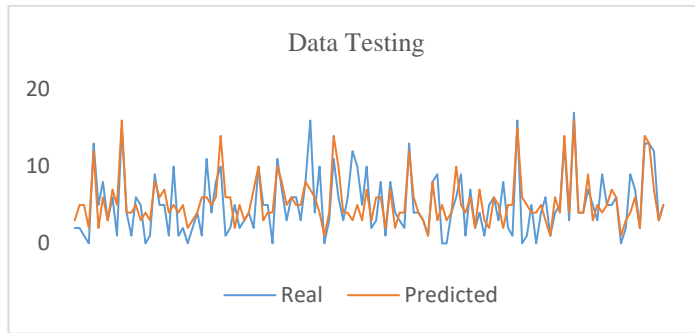


Figure 8. The real and prediction line chart of data training from 2017-2018.

Table 5 Two-samples t-test for Data Testing

	Real	Predicted
Mean	5.126984	5.428571
Variance	16.01575	10.10286
Observations	126	126
Hypothesized Mean Difference	0	
df	238	
t Stat	-0.6624	
P(T<=t) one-tail	0.254176	
t Critical one-tail	1.651281	
P(T<=t) two-tail	0.508353	
t Critical two-tail	1.969982	

3.4 Web implementation

This modeled is implemented in a website base to help the “Dinas Kesehatan Surabaya” (The Surabaya Public Health Department) monitoring the dengue fever outbreak. From this website, users can see DF spreading data for each district in Surabaya in the selected year (Figure 9). Data on the number of victims in each sub-district will be displayed in red, yellow and green, with red representing the largest number of victims and green representing the smallest number of victims. Users can specify the upper limit of each color representative. The application will then automatically determine the color gradation based on the input, so that the user can see number of victims in each sub-district that representing in color information. The legend from this gradation color information will display next to the map. User also could choose and see detail information from each sub-district and number of DF victims.

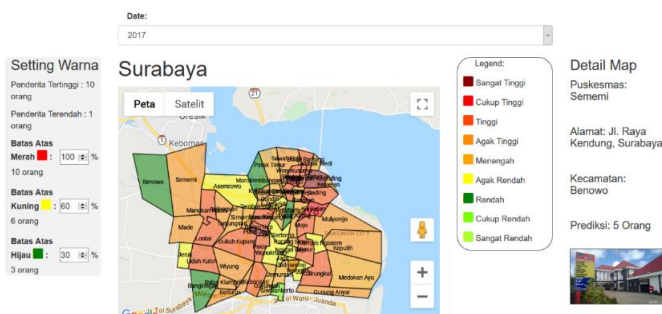


Figure 9. The web design for implemented model

Users can also see details of the number of patients in each sub-district and compare the movement of the number of patients in 3 years presented in tabular form (Figure 10). This data also can be viewed in graphical form (Figure 11).

This visual information will provide more informative information to help The Surabaya Public Health Department monitoring and prevent the dengue fever outbreak for each sub-district.

Table Summary

Puskesmas	Jumlah Penderita 2012	Jumlah Penderita 2013	Jumlah Penderita 2014
Asemrowo	16 orang	22 orang	5 orang
Balas Klumprik	8 orang	20 orang	4 orang
Balongsari	32 orang	46 orang	11 orang
Bangkingan	8 orang	6 orang	7 orang
Banyu Urip	29 orang	48 orang	19 orang
Benowo	23 orang	35 orang	6 orang
Bulak Banteng	12 orang	7 orang	5 orang
Dr. Soetomo	14 orang	43 orang	6 orang
Dukuh Kumpang	25 orang	66 orang	25 orang
Dupak	3 orang	8 orang	6 orang

Figure 10. Summarize comparizon spreading DF disease data for each sub-district in Surabaya

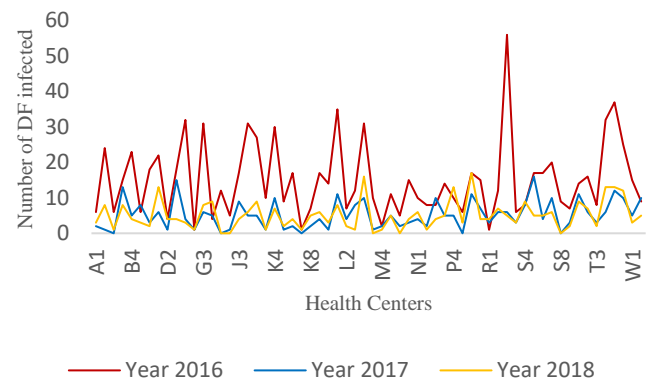


Figure 11. Summarize comparizon spreading DF disease data for each sub-district in Surabaya

4. Conclusion

In this paper we proposed spatial multi-layer perceptron (spatial MLP) model for predicting dengue fever in Surabaya. The model can capture the data pattern. Additionally, the model is implemented in the web-based database. The Surabaya Public Health Department (Dinas Kesehatan Surabaya) can input the data and predict the outbreak online. However, right now in some regions the predictions are not performed well, especially when that region has zero value. The zeros will be predicted as the mean values it's neighborhood. In the next research, we will expand the model into spatial-temporal multi-layer perceptron (spatial-temporal MLP) model, which can capture data dependencies not only spatially, but also temporally.

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