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Optimization of Naïve Bayes uses the genetic algorithm for classification data

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Abstract. Classification is one of the statistical methods to classify data systematically. However, if there is a large amount of data and various features, it often results in low accuracy. For this reason, methods are needed that can handle the data with various types. One method that can handle this problem is Naïve Bayes. Naïve Bayes is one of the methods used for classification data. This method requires a stage of selection of independent variables in increasing the accuracy of the model from Naïve Bayes. So we need an excellent method to fix these deficiencies uses a Genetic Algorithm (GA). Genetic algorithm is one of the metaheuristic methods used in optimization techniques. The data used are septic tank data in East Surabaya with eleven independent for classification data. The result of classification accuracy using Naïve Bayes is 72.7%. When Naive Bayes was used with a genetic algorithm, the classification accuracy was increased is 90.9%

1. Introduction

Classification is a statistical method for group or classifies the data systematically arranged. The classification problems arise when several rules that consist of one or more categories can not be identified immediately but must use a size. In statistics, several classification methods are used to per arm data classification, such as discriminant analysis, logistic regression, Naïve Bayes, etc. [1].

Naïve Bayes is a simple probabilistic classification that calculates probabilities by summing the gequency and the combined value of a given dataset. In its application, Naïve Bayes is a method that requires a small number of training data to determine the estimated parameters required in the process classification. Naïve Bayes often works better in most real-world situations complex than expected [2]. The Naive Bayes method required stages of selecting variables to improve the model's accuracy in explaining the data. So that needed another method in variable selection one of them is Genetic Algorithm (GA). Genetic Algorithm (GA) is one of the metaheuristic methods used in optimization techniques. According to Haupt and Haupt [3], a Genetic Algorithm is an optimization technique based on genetic principles and natural selection. In the Genetic Algorithm, the population comprises many individuals who develop according to specific selection rules by maximizing fitness. According to



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Sivanandam and Deepa [4], the advantages possessed by Genetic Algorithm compared to other methods are very suitable for solving global problems optimum, easy to change, or flexible to implement on various problems more expansive solution space. According to Xu and Zhang [5], in their research stated that the selection of variables using the genetic algorithm approach is better than the method of selection of backward elimination, forward selection, and stepwise method for multiple regression. According to research [6-10], naïve Bayes with genetic algorithms is a good combination of implementing applications. According to this study [11,12], the genetic algorithm is used to optimize logistic regression, and the results are better than without the genetic algorithm.

This study aims to know the optimization of naïve bayes using a genetic algorithm for classification data based on the above description. The data used to see the effect is the data on domestic waste results in the Surabaya area, where this data is classified as data [13].

2. Methods

2.1. Naïve Baves

Naïve Bayes is a simple odds classification that calculates multiple odds by adding up the frequency and value combinations from a specific data set. Naive Bayes the Bayes theorem and assumes all independent attributes are assigned to class variables[14]. Naïve Bayes is a classification with probability and statistical methods [6].

The advantage of using Bayes is that this method only requires a small amount of training data to determine the classification process's estimated parameters. Naïve Bayes often works far better in most complex real-world situations than expected [2].

| The equation from the Bayes theorem is: | |
|---|-----|
| $P(H \mid X) = \frac{P(X \mid H).P(H)}{P(X \mid H).P(H)}$ | (1) |

 $\frac{P(H \mid X)}{P(X)} = \frac{P(X)}{P(X)}$ To explain the Naïve Bayes method, it is essential to know that the classification process requires several

clues to determine what class is suitable for the analyzed sample. Therefore, the Naïve Bayes method above is adjusted as follows:

$$\underline{P(C \mid F1...Fn)} = \frac{\underline{P(C)P(F1...Fn \mid C)}}{\underline{P(F1...Fn)}}$$
(2)

The variable C represents the class, while the variable F1 \cdots Fn presents the characteristic instructions needed to do the classification. Further elaboration of the Bayes formula is carried out by describin $(C = F1, \cdots, Fn)$ using the multiplication rule. A very high independence assumption (naïve) is used, that each of the instructions (F1, F2, \cdots , Fn) is independent of each other. With these assumptions, the following similarities apply:

$$P(F_i | F_j) = \frac{P(F_i \cap F_j)}{P(F_j)} = \frac{P(F_i)P(F_j)}{P(F_j)} = P(F_i), \text{ for } i \neq j,$$
(3)

🛜 that,

$$P(F_i \mid C, F_j) = P(F_i \mid C)$$
(4)

The equation above is a model of the Naïve Bayes theorem, which will then be used in the classification process. For classification with continuous data, the Gauss Density formula is used:

$$P(X_{i} = x_{i} | Y = y_{j}) = \frac{1}{\sqrt{2\pi\sigma_{ij}}} e^{\frac{(X_{i} - A_{ij})^{2}}{2\sigma^{2}_{ij}}}$$
(5)

2.2₂ Genetic Algorithm (GA)

A genetic algorithm is an optimization technique based on genetic principles and natural selection. Genetic algorithms are formed from many individuals who develop according to specific selection rules by maximizing fitness [15]. This algorithm is also used to get the optimum global value by repeating or iterating the Darwinian concept of evolution.

According to Trevino and Falcian [16], there are 7 stages for carrying out genetic algorithms, namely:

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|---------------------------------------|---------------------------|-------------------------------------|
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- 1. It was forming an initial population consisting of several chromosomes which contain chromosome genes in a genetic algorithm that is used to indicate candidates for a group of genes that can be used as solutions to problems. Genes in genetic algorithms contain variables that want to be optimized. In this study, genes contain independent variables.
- 2. Each chromosome in the population is evaluated for its ability by using the fitness function. In this study, the fitness function is in the form of a misclassification.
- 3. When a chromosome has a more optimum fitness value than its initial value, then the chromosome is stopped, but if not, then the analysis stage is continued to stage 4. The smallest fitness value chosen as a solution to this study's problem because the fitness function used is the level of misclassification.
- 4. It was choosing chromosomes with optimum fitness values that parents use.
- 5. They were combining genetic information in parent's replication through crossing over. Two parents are randomly selected and used to form two new chromosomes.
- 6. We are doing mutations to introduce new gene elements to chromosomes randomly.
- 7. Stages are repeated from step 2 to the chromosomes that provide the most optimal fitness value or have reached convergence

2.3. Evaluation of Classification Method Performance

Actual data and predictive data from the classification model are presented using a cross-tabulation (Confusion matrix), which contains information about the actual data class represented on the matrix row and the predictive data class in column [17]. Classification accuracy can be seen in Table 1.

| | Table 1. | Classific | ation Tabl | le | |
|--|--------------------------|-------------------------|--------------------------|--|------------------------------|
| | Actual | Pred | iction | | |
| | Actual | Positif | Negatif | | |
| | Positif | TP | FN | | |
| | Negatif | FP | TN | | |
| where, | | | | | |
| $accuration = \frac{TN + TP}{TN + TP + FN + FP}$ | | | | _ | (6) |
| Besides accuracy, classification per values. Sensitivity is the accuracy of | rformance the positiv | can also ve class, v | be assesse while spec | ed based on sensitivity and ficity is the accuracy of t | l specificity he negative |
| class. The formula for sensitivity and | | | | | |
| $Sensitivity = \frac{TP}{(TP + FN)} \times 100\%$ | | | | | (7) |
| $Specificity = \frac{TN}{(TN + FP)} \times 100\%$ | | | | | (8) |

Also, performance evaluation of the classification model can be done using the G-mean. G-mean is the average geometric sensitivity and specificity. If all positive classes cannot be predicted, the G-mean will be zero, so a classification algorithm is expected to achieve a high G-mean value [11] $G - Mean = \sqrt{Sensitivity \times Specificity}$ (9)

3. Results and Discussion

3.1. Analysis Naïve Bayes

Naïve Bayes methods of use in classifying a ratio of 90% for the training data with 10 % for the data testing. Then we count the prior probability of Y in Table 2.

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Tabel 2. Prior Probability of Naïve Bayes

| У | · |
|-------|-------|
| 0 | 1 |
| 0,308 | 0,692 |

And then, calculate the odds of each of the marginal opportunities for each variable X against the variable Y. The following is given. The results of the opportunities for each variable are summarized in Table 3.

| T | a b | r · 1 | 1 . | 1.111. |
|----------|------------|----------|-------|--------|
| able | 5. IV | larginal | proba | bility |
| | | | | |

| | | | | | | | X6 | | |
|----------|-------|----------------|-------|-------|----|-------|-------|-------|-----|
| у | х | 1 | | | У | 0 | 1 | 2 | |
| <i>y</i> | 0 | 1 | | | 0 | 0,500 | 0,214 | 0,214 | 0,0 |
| 0 | 0,750 | 0,250 | | | 1 | 0,492 | 0,318 | 0,191 | 0,0 |
| 1 | 0,619 | 0,381 | | | | х | 7 | | |
| v | | х | 2 | | У | 0 | 1 | | |
| У | 0 | 1 | 2 | 3 | 0 | 0,000 | 0,286 | | |
| 0 | 0,179 | 0,286 | 0,357 | 0,179 | 1 | 0.079 | 0,143 | | |
| 1 | 0,286 | 0,238 | 0,429 | 0,048 | | х | 8 | | |
| | | х | 3 | | У | 0 | 1 | | |
| У | 0 | 1 | 2 | 3 | 0 | 0,107 | 0,893 | | |
| 0 | 0,214 | 0,321 | 0,250 | 0,214 | 1 | 0,064 | 0,937 | | |
| 1 | 0,016 | 0,254 | 0,333 | 0,397 | | | X9 | | |
| | | \mathbf{X}_4 | | | У | 0 | 1 | 2 | |
| У | 0 | 1 | | | 0 | 0,357 | 0,036 | 0,036 | 0,: |
| 0 | 0,214 | 0,78 | 6 | | 1 | 0,302 | 0,206 | 0,079 | 0,4 |
| 1 | 0,508 | 0,49 | 2 | | ., | X | 10 | | |
| v | | x5 | | | У | 0 | 1 | | |
| У | 0 | 1 | | | 0 | 0,321 | 0,679 | | |
| 0 | 0,607 | 0,39 | 3 | | 1 | 0,191 | 0,809 | | |
| 1 | 0,873 | 0,12 | 7 | | | X | 11 | | |
| | | | | | У | 0 | 1 | | |
| | | | | | 0 | 0,607 | 0,393 | | |
| | | | | | 1 | 0,318 | 0,683 | | |

After calculating the prior and marginal opportunities, then the prediction results obtained from the Naïve Bayes probability model are used to compare the probability of testing data 10% of 102 observations, 11 observations, and are written in Table 4, as follows:

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| Table 4. | Prediction y |
|----------|--------------|
| У | ŷ |
| 1 | 1 |
| 1 | 0 |
| 0 | 1 |
| 1 | 1 |
| 0 | 0 |
| 0 | 1 |
| 0 | 0 |
| 1 | 1 |
| 1 | 1 |
| 1 | 1 |
| 1 | 1 |

from the results of the predictions in Table 4, then the actual data is compared, and the Naïve Bayes contingency table is formed in Table 5, as follows:

Table 5. Table of contingencyNaïve Bayes

| Real | Predi | ction |
|------|-------|-------|
| Keai | 0 | 1 |
| 0 | 2 | 1 |
| 1 | 2 | 6 |

From Table 5, the accuracy of the Naïve Bayes classification will be calculated summarized in Table 6, as follows:

| Table 6. Classification Accuracy of Naïve Bayes | | | | | | | | |
|---|-------|-------------|-------------|--------|--|--|--|--|
| Akurasi | APER | Specificity | Sensitivity | G-Mean | | | | |
| 72.7% | 27.3% | 0.9 | 0.5 | 0.7 | | | | |

3.2. Selection analysis of GA - Naïve Bayes

The genetic algorithm stage will stop if the minimum fitness value converges from the previous and next generation. The results of the genetic algorithm with the Naïve Bayes model can be seen in Table 7 as follows:

| Chromosome | | | | | | | Fitness |
|------------|----------------|----------------|----------------|----------------|-------------------|-------------------|---------|
| Chromosome | \mathbf{x}_1 | \mathbf{X}_2 | \mathbf{X}_3 | X 9 | \mathbf{x}_{10} | \mathbf{x}_{11} | Values |
| 1 | 0 | 1 | 0 | 1 | 1 | 1 | 0,134 |
| 2 | 0 | 0 | 1 | 1 | 1 | 1 | 0,134 |
| | | | | | | | |
| | | | | | | | |
| | | | | | | | |
| 99 | 0 | 0 | 1 | 1 | 1 | 1 | 0,134 |

Table 7. There are 100 Chromosome Results and Fitness Values

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| Chromosome | | | | | | | Fitness |
|------------|----------------|----------------|------------|----------------|-------------------|-------------|---------|
| Chromosome | \mathbf{x}_1 | \mathbf{x}_2 | X 3 | X 9 | \mathbf{X}_{10} | X 11 | Values |
| 100 | 0 | 0 | 1 | 1 | 1 | 1 | 0,134 |

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From Table 7, the results of the Genetic Algorithm selection above obtained 5 selected variables, namely x_{6} , x_{7} , x_{9} , and x_{11} contained on the first chromosome with a fitness value of 0.134. The following Table 8 will summarize the results of the classification accuracy.

Table 8. The Classification Accuracy of Naive Bayes with Genetic Algorithm

| | | | | | G- |
|---|----------|------|-------------|-------------|------|
| _ | Accuracy | APER | Specificity | Sensitivity | Mean |
| | 90,9% | 9,1% | 1 | 0,8 | 0,9 |

3.3. Comparison of Classification accuracy

From the results obtained, the comparison of the methods used between Naïve Bayes with Genetic Algorithms - Naïve Bayes in Table 9, as follows:

| Table 9. Comparison of Classification accuracy | | | | | | | |
|--|---|--|--------|-----|-----|-----|--|
| Method | Selection | The measure of classification accuracy | | | | | |
| Method | Variable | 1 | 2 | 3 | 4 | 5 | |
| Naïve Bayes | x ₁ , x ₂ ,, x ₁₁ | 72,7 % | 27,3 % | 0,9 | 0,5 | 0,7 | |
| GA - Naïve Bayes | x ₆ , x ₇ , x ₉ , x ₁₁ | 90,9 % | 9,1 % | 1 | 0,8 | 0,9 | |
| Information : | | | | | | | |
| 1: Accuracy | | | | | | | |
| 2: APER | | | | | | | |
| 3: Specificity | | | | | | | |
| 4: Sensitivity | | | | | | | |
| 5: G-Mean | | | | | | | |

Based on Table 9, it can be seen that naïve Bayes with a genetic algorithm has a better classification accuracy than the results without a genetic algorithm. Then the variables selected smaller than the total generated by the naïve Bayes only.

4. Conclusion

The results of data analysis result from classification accuracy Genetic Algorithm - Naïve Bayes has an accuracy of 90,91% dan APER is 9,09%. But the result of variable selection Naïve Bayes is less than is 4 variable.

References

- Salim A 2017 Pengoptimalan Naive Bayes Dan Regresi Logistik Menggunakan Algoritma Genetika Untuk Data Klasifikasi. Unpublished.
- [2] Patterkari S A and Parveen A 2012 Internastional J. Adv. Comput. Math. Sci. 3 290 294.
- [3] Haupt R L and Haupt S E 2003 Practical Genetic Algorithms (Hoboken, NJ, USA: John
- Wiley & Sons, Inc).
 [4] Sivanandam S N and Deepa S N 2008 *Genetic Algorithms in Introduction to Genetic Algorithms* (Berlin, Heidelberg: Springer Berlin Heidelberg, 15 37)
- [5] Xu L and Zhang W J 2001 Anal. Chim. Acta. 446 475.

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- [6] Bustami 2013 J. Penelitian Teknik Inform. 3 127 146
- [7] Buani D C P 2016 EVOLUSI J. Sains dan Manaj 4 1
- [8] Choubey D K, Paul S and Kumar S 2017 Classification of Pima indian diabetes dataset using naive bayes with genetic algorithm as an attribute selection.
- [9] Govindarajan M 2013 Sentiment Analysis of Movie Reviews using Hybrid Method of Naive Bayes and Genetic Algorithm.
- [10] Busono S 2020 J. Ilm. Teknol. Inf. Asia. 14 31
- [11] Wardhani R P S, Sudarno S and Maruddani D A I 2019 J. Gaussian 8 506 51
- [12] Salim A and Alfian M R 2019 J. Teknol. Inf. dan Terap. 6 2580 2291
- [13] Kusumawati Y 2013 Pemodelan Faktor-Faktor Yang Mempengaruhi Rumah Tangga Membuang Limbah Domestik Menggunakan Regresi Logistik Dan Algoritma Genetika. Unpublished.
- [14] Patil T R and Shereker M S 2013 Int. J. Comput. Sci. Appl. 6 256 261
- [15] Han J, Kamber M and Pei J 2006 Data mining: concepts and techniques, Second ed. (San Francisco)
- [16] Trevino V and Falciani F 2006 Bioinformatics 22 1154 1156
- [17] Miroslav K and Matwin S 1965 Br. J. Psychiatry. 111 1009 10

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