Estimation of biomass of forage sorghum (sorghum bicolor) Cv. Samurai-2 using support vector regression

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Article Info ABSTRACT

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Keywords:

Estimated production Forage sorghum Machine learning Sorghum bicolor Support vector regression One alternative to improve feed quality is to combine the main feed with forages which are more economical in cost but contain high protein sources, such as sorghum. Production estimation is essential because it will determine the sustainability of the feed. This study aimed to estimate the amount of sorghum production using support vector regression (SVR). Several stages of this research are collecting data, preprocessing, modelling, and evaluation. The dataset used and the input for this SVR algorithm model is field observation data. The kernels used in the SVR algorithm modelling are linear, Polynomial, and RBF. Sorghum production estimation using SVR has a performance evaluation value that refers to the root mean square error (RMSE). The result of this research is that the model obtained from the SVR algorithm can estimate sorghum production with performance evaluation values using R2, mean absolute error (MAE), mean absolute percentage error (MAPE), and RMSE. The best results on the Polynomial kernel are R2=0.7841, MAE=0.0681, MAPE=0.46641, and RMSE=0.1006. This study shows that the classification model obtained from the SVR algorithm with Kernel Polynomial is the best model for estimating sorghum production.

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1. INTRODUCTION

In the livestock industry, a significant concern is the availability of land, livestock and feed. The main feed for the livestock industry is grass, but the grass has high fibre and low protein content for livestock. The grass needs to be mixed with concentrates such as dregs, corn, and other similar foods to add nutritional value to the feed. It will cause production costs to increase. Abdullah and Suharlina [1], an alternative source of high forage protein but at an economical cost is to combine the main feed with types of legumes such as sorghum. Sorghum has been introduced and cultivated in Indonesia, particularly in dry and marginal areas [2], and it is a universal multipurpose crop for food, fodder, and potential biofuel feedstock [3].

The problem in this study is that to estimate the production of feed biomass for ruminants is sufficient or more or less as a feed production material to combine forage with the main feed. Therefore it is necessary to estimate the biomass of sorghum yields. In the industrial era 4.0, information technology has become necessary in various fields, including agriculture and animal husbandry. One of the uses of this information technology is the application of machine learning algorithm models, namely computer programming to optimize performance using history data [4]. Liakos *et al.* [5], in general, the machine learning methodology involves a learning process to learn from training data in carrying out tasks. It is stated by Ghosal *et al.* [6] in the national strategy for artificial intelligence in Indonesia that increasing forecasting accuracy with machine learning will help farmers plan agricultural cycle activities. Several previous studies by BPPT [7] and Masjedi et al. [8] used machine learning algorithms for research on sorghum plants. Several studies related to predictions in agriculture and animal husbandry are wheat yield prediction [9], crop disease prediction [10] and [11]. One widely used algorithm for forecasting or predicting target values is the support vector machine (SVM), for example, in the medical field [12], analogue circuit [13], education [14], face recognition [15] and also in the agriculture [16]. For solving the SVM regression case, it is modified to the support vector regression (SVR) algorithm [17]. SVR aims to find a hyperplane to predict the training data set and the optimal value of the parameters obtained through the GridSearch method. The grid search method tests a model to find the error value in the classification [18]. The parameters determined by the optimal value are epsilon (ϵ), cost (C), and gamma (γ). In the SVR method, several choices of kernel functions can be used, such as linear, Gaussian, Polynomial, and several other kernels. SVR builds a hyperplane in high dimensional space in linear or nonlinear data and can overcome overfitting [19]. Several studies using SVR in agriculture and animal husbandry, such as soil erosion susceptibility prediction [20], predicting forage quality of warm-season legumes [21], crop model of rice production [22], Water stress detection [23] and also Rumex and Urtica detection in grasslands [24]. In this study, it is hoped that the machine learning SVR algorithm models will be widely used to predict or estimate the amount of sorghum biomass production. This study will look for the best kernel function of the three kernels and the best parameters using the GridSearch method from the SVR model to estimate the biomass of animal feed sorghum production.

2. METHOD

This research methodology has five stages: a preliminary study, data collection, preprocessing, modelling, and model evaluation. Each step in this research flow cycle diagram is as shown in the research flow chart Figure 1. The explanation of each stage of this research carried out is as:

- Data collection: This study uses the sample data taken from direct observations in the sorghum field in the sorghum bicolor block cv. Samurai-2. The research area is Jonggol Animal Science Teaching and Research Unit (JASTRU), Singasari Village, Jonggol District, Bogor City. The dataset was taken at harvest time on March 8, 2021, with as many as 88 plant data with attributes shown in Table 1 and sample data from the field shown in Table 2.

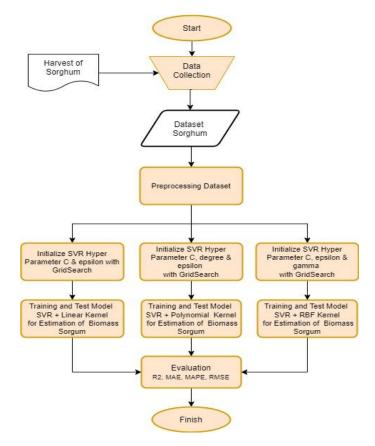


Figure 1. Research steps

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Mo

| No | Latitude | Longitude | Stem | Stem | Leaves | Seed | Seed | Biomass |
|--------|-------------|--------------|--------|----------|--------|--------|-------|---------|
| sample | | | height | diameter | | height | width | |
| 6 | -6.46875233 | 107.01083981 | 170 | 14 | 9 | 19 | 4 | 204 |
| 12 | -6.46885800 | 107.01121700 | 224 | 25 | 15 | 24 | 5 | 677 |
| 18 | -6.46867571 | 107.01117843 | 125 | 11 | 10 | 22 | 4 | 147 |
| 27 | -6.46846183 | 107.01120257 | 219 | 16 | 10 | 27 | 7 | 291 |
| 38 | -6.46855311 | 107.01108322 | 219 | 23 | 11 | 32 | 9 | 574 |
| 39 | -6.46838988 | 107.01111775 | 217 | 20 | 12 | 25 | 5 | 381 |
| 42 | -6.46840753 | 107.01109931 | 177 | 19 | 11 | 19 | 4 | 341 |
| 43 | -6.46847116 | 107.01105975 | 219 | 20 | 10 | 28 | 6 | 480 |
| 44 | -6.46856444 | 107.01105069 | 150 | 15 | 10 | 18 | 3 | 156 |
| 47 | -6.46861375 | 107.01098096 | 232 | 19 | 11 | 29 | 6 | 306 |
| 48 | -6.46849082 | 107.01104090 | 234 | 22 | 12 | 30 | 8 | 554 |
| 50 | -6.46826595 | 107.01107886 | 164 | 15 | 9 | 26 | 5 | 231 |
| 52 | -6.46839654 | 107.01091088 | 215 | 16 | 13 | 26 | 5 | 271 |
| | | | | | | | | |

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Table 2. Attribute dataset sorghum of bicolor cv Samurai-2

| - | NO | Attribute | Description |
|---|----|---------------|--|
| | 1 | NO_SAMPLE | No samples, A and B indicate growing two plant stems in one plant location |
| | 2 | LATITUDE | Latitude |
| | 3 | LONGITUDE | Longitude |
| | 4 | STEM_HEIGHT | Plant stem height in meters |
| | 5 | STEM_DIAMETER | Plant stem diameter in millimeters |
| | 6 | LEAVE | Number of leaves |
| | 7 | SEED_HEIGHT | Sorghum seed length in centimeters |
| | 8 | SEED_WIDTH | Sorghum seed width in centimeters |
| | 9 | BIOMASS | Overall weight in grams (target class) |
| | | | |

- Preprocess: The attributes of the dataset from field obtained recording as shown in Table 1. The target attribute is WEIGHT, which is the plant's total weight (gram). All selected features are normalized between 0 and 1 before input into the SVR algorithm model.
- Modeling: At this stage, the preprocessed dataset will be input into each SVR algorithm with a different kernel function, namely the SVR algorithm with a linear kernel function, the SVR algorithm with a gaussian kernel function, and the SVR algorithm with a polynomial kernel function. Several parameter values are identified to produce the best results from each kernel function in the modelling and evaluation process. The model validation process uses k-fold cross validation.
- Evaluation: This stage is the output of the modelling process using the validation process. That is to
 compare actual and predicted values with a low error rate from every kernel function of the SVR. That
 model selected will be recommended for the prediction of sorghum production. Validation of all kernel
 functions is using R-Squared, mean absolute error (MAE), mean absolute percentage error (MAPE), and
 root mean squared error (RMSE). The achievement of this final stage indicator is that the resulting model
 has a minimum error value.

3. RESULTS AND DISCUSSION

Data correlation analysis on dataset attributes from collecting selected sorghum research sample data at harvest time, namely STEM_HEIGHT, STEM_DIAMETER, LEAVE, SEED_HEIGHT, SEED_WIDTH, and BIOMASS, there is very high correlation data. Also, data with weak correlation and visually correlation data between these attributes are shown in Figure 2. The highest data correlation was on the seed or panicle width attribute (seed_width) with the seed or panicle length attribute (seed_height), which was 0.92. The number of leaves (leaves) with a length is a weak correlation of a collection of seeds or panicles (seed_height) of 0.35. The preprocessing, modelling, and evaluation stages are carried out using the Python 3.7 programming language. Python using the skit-learn library and several main class packages for vector regression and cross-validation.

In the preprocessing stage, scalar normalization was carried out to convert the value proportionally in each attribute between 0 to 1. The preprocessed dataset is the input of the SVR algorithm with a vector regression package at the modelling stage. The output of the modelling process is validated, and the results are evaluated at the evaluation stage. The standard value of a suitable validation parameter in using the cross-validation method specified to determine accuracy is ten times [25]. The SVR model will construct a hyperplane in high dimensional space in the nonlinear data shown in (1).

 $(x_i) = (w.x_i) + b$

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Where:

f(x) = predictive value

x = dataw = weight

 $b = \text{bias value}, (\text{also represented by}\lambda)$

Estimation for the coefficients w and b through the risk function and with |w| as the normalization of the function to minimize it to produce a function close to flat, E_{ε} is an ε -insensitive loss function. The coefficient C value is defined by the user (trade-off) between the thin distance of the f function and the value above the upper limit deviation, which can still be tolerated as shown in (2) [26].

$$R(f(x_i)) = \frac{1}{2} \| w \|^2 + C \sum_{i=1}^n E_{\varepsilon} (y_i - f(x_i))$$
(2)

$$f(\mathbf{x}_i) = \sum_{i \in R} (\alpha_i - \alpha_i^*) K(\mathbf{x}_i, \mathbf{x}) + b$$
(3)

Smola and Schölkopf [27] from (2), determining the parameters w and b become an optimization problem using the Lagrangian. The final equation for determining predictions with SVR is shown in (3), where are the Lagrange multiplier and the selected kernel function. Several kernel functions to handle nonlinear data cases are often used in SVR models, such as linear kernel in (4), polynomial kernel in (5), and Gaussian or Gaussian radial basis function (RBF) kernel in (6).

$$K(x_i, x) = (x, x') \tag{4}$$

$$K(x_i, x) = (\gamma < x, x' > +r)^d \tag{5}$$

Where d is the degree parameter and r is the coefficient.

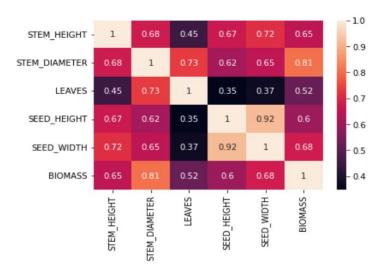
$$K(x_i, x) = \exp(-\gamma || x - x' ||)2$$
(6)

Where is the gamma parameter must be greater than 0.

Figure 2. Dataset correlation

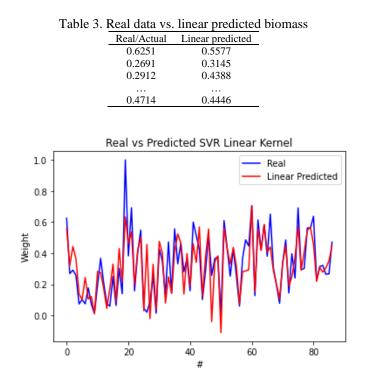
The complete grid search process time is very long. Therefore, Hsu and Lin [18] recommends performing the grid search through the loose grid stage for selecting C and values, then proceeding with the finer grid stage to get a value around the C value that has been obtained with the lowest error value previously. The search for parameters C and epsilon on the SVR model in each kernel is done using GridSearch with a combination of parameter values tested, namely C = 0.01, 0.1, 1, 100, 1000 and epsilon parameters = 0.0001, 0.0005, 0.001, 0.005, 0.1, 0.05, 0.1, 0.5, 1, 5, 10.

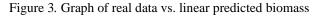
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(1)

The modelling process with a linear kernel obtained the best parameters for C and epsilon through the GridSearch method were C=1 and epsilon=0.0005, resulting in predictive values compared to real or actual values, as the example shown in Table 3, and graphically visualization is shown in Figure 3. Then the modelling process with a polynomial kernel obtained the best parameters for C, degree, and epsilon through the GridSearch method were C=1, degree=2, and epsilon=0.001, which resulted in predictive values compared to real or actual values as the example shown in Table 4, and graphically visualization is shown in Figure 4.





| | cui uutu vo. | porynomial predicte | | |
|---|--------------|----------------------|--|--|
| | Real/Actual | Polynomial Predicted | | |
| | 0,6251 | 0,5725 | | |
| | 0,2691 | 0,2869 | | |
| | 0,2912 | 0,4175 | | |
| | | | | |
| _ | 0,4714 | 0,4734 | | |
| | | | | |

| Table 4. F | Real data vs | s. polynomial | predicted biomass |
|------------|---------------|---------------|-------------------|
| | Da -1/A atra- | D-1 | Due d'ate d |

Furthermore, the modelling process with the RBF kernel obtained the best parameters for C, epsilon, and gamma through the GridSearch method were C=100, epsilon=0.01, and gamma=0.1, which resulted in prediction values compared to real or actual values as the example shown in Table 5 and graphically visualization is shown in Figure 5.

| Table 5. Rea | l data vs. lin | ear Gaussian F | RBF biomass |
|--------------|----------------|----------------------|-------------|
| | Real/Actual | RBF Predicted | |
| | 0,6251 | 0,5813 | |
| | 0,2691 | 0,2846 | |
| | 0,2912 | 0,3591 | |
| | | | |
| | 0,4714 | 0,4767 | |

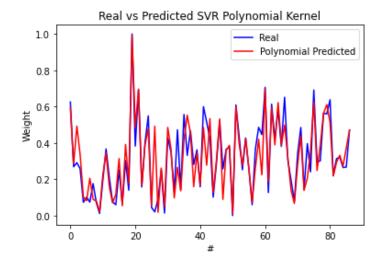


Figure 4. Graph of real data vs. polynomial predicted biomass

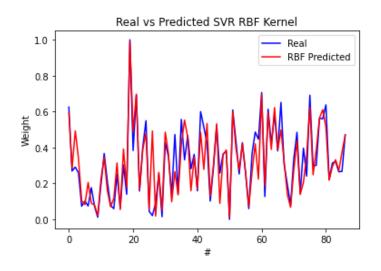


Figure 5. Graph real data vs Gaussian RBF predicted biomass

Measurement of accuracy and measurement of the error value between the predicted value and the real or actual value at the evaluation stage for each model result for each kernel uses the R-squared accuracy measurement method in (7) [28].

$$R^2 = 1 - \frac{SSER}{SSEM}$$
(7)

Meanwhile, for measuring the error value between the predicted value and the real or actual value at the evaluation stage for each model result, each kernel uses the MAE error measurement method [29] in (8), MAPE in (9), and RMSE in n (10) [28].

$$MAE = \frac{1}{n} \sum_{1}^{n} (y^{\wedge} - y) \tag{8}$$

MAPE =
$$\frac{1}{n} \sum_{1}^{n} (y^{\circ} - y) x 100$$
 (9)

$$\text{RMSE} = \sqrt{\frac{1}{n} \sum_{1}^{n} (y^{\wedge} - y)^2}$$
(10)

The predicted value and the real or actual value of the modelling results from each kernel are entered into the MAE, MAPE, and RMSE measurement equations to produce error measurement values, shown in Table 6.

| Table 6. Metric comparison result | | | | |
|-----------------------------------|---------------|-------------------|---------------------|--|
| | Linear Kernel | Polynomial Kernel | Gaussian RBF Kernel | |
| R-squared | 0.6861 | 0.7841 | 0.7751 | |
| MAE | 0.0836 | 0.0681 | 0.0715 | |
| RMSE | 0.1173 | 0.1006 | 0.1055 | |
| MAPE | 55.1891 | 46.641 | 48.7648 | |

SVR with Polynomial Kernel has the smallest MAE, MAPE, and RMSE error measurements compared to the linear kernel and the Gaussian RBF kernel. Likewise, with the R-squared value of the polynomial kernel compared to the linear kernel and Gaussian kernel, the polynomial kernel has the largest R-squared value, so based on the research that has been done that the Polynomial Kernel is a good Kernel on the SVR method for estimating the production of forage sorghum (sorghum bicolor) cv. Samurai-2.

4. CONCLUSION

In this study, to obtain an SVR model for estimating the biomass of animal feed sorghum, we tried the SVR model with a combination of linear kernel, polynomial kernel, and gaussian RBF kernel. Each of our kernels looks for the best parameters using the GridSearch method function. The result is that the SVR model using the Polynomial Kernel kernel function with parameters C=1, degree=2, and epsilon=0.001 has the lowest error value and the highest coefficient of determination. Thus, SVR with a polynomial kernel function can be recommended to estimate the biomass of sorghum bicolor cv Samurai-2. The prospect of developing research results and implementing further research in the future can use other kernels such as Splines, B-Splines, and others.

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Estimation of biomass of forage sorghum (sorghum bicolor) Cv. Samurai-2 ... (Kahfi Heryandi Suradiradja)



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