

Covid-19 Patient Care Status Using Random Forest and Support Vector Machine Methods

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Abstract—Covid-19 disease spread throughout the world and caused death globally. Accordingly researchers use many research methods to control the Covid-19 pandemic, one of which is getting more attention is the use of prediction methods. The problem in medical action is not easy to determine the treatment status. So it is not surprising that previous researchers emphasized that the most frequent mistakes were due to inaccuracies in medical decisions. Therefore, this study proposes a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not only because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analytics for better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. So, in essence, it is not easy to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of the field of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. In fact, machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [19][20][21]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [22]. For this reason, or that is why this study proposes a machine learning system model for decision-making solutions (predictions) of the treatment status of Covid-19 patients, both inpatient and self-isolation using data mining methods.

Two things machine learning can predict, are classification to predict class membership and regression to show numerical values [12]. While data mining is actually part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in the field of prediction [23]. The

embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting the new data with high accuracy [22].

There are many Data Mining methods, including K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out in order to be able to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [24] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [25] [26], as well as SVM algorithm has a widely known method used for classification [26].

However, it is essential to know the level of accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study, whether the patient should be hospitalized or self-isolated. Therefore, this study also conducted a further test of the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. Testing the accuracy of predicting the treatment status of Covid-19 patients is carried out on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some of the related works from the latest scientific articles compared with the work in this research article conducted.

- Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [24]. This previous research is different from the research in the article on the research objectives and the object

under study.

- Celestine Iwendi et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [27]. The difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM methods. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.
- Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [28]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, the research in this article predicts the treatment status of Covid-19 patients.
- Based on patient clinical data, Boran Hao et al. (2020) developed a model to predict the severity of pneumonia and the level of care required by Covid-19 patients using statistical methods.[29]. This previous research differs in the research purpose and method compared to the research in this article.
- Anthony Anggrawan et al. (2021) implemented machine learning to diagnose drug users and types of drug users using Forward Chaining and Certainty Factor methods [22]. Mean while, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.
- Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future by using the Poisson distribution and the gamma distribution [30]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article.
- Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. The research in this article is different from previous research; the difference lies in the research objectives and the research methods used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. The research in this article is not using the DT method but uses the RF and SVM methods.
- Soham Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [31]. This previous study differed in its research objectives from the research in this article. The

previous research predicts patients suffering from Covid-19 or not using the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods.

- Ankit Mehrotraa and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [32]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from the research in this article because this article is an experimental study, not a literature review research.
- Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods used compared to the research in this article. The difference lies in the fact that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].
- Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [33]. However, the previous research differs in the purpose and the research method compared to the research in this article.
- Vadim Demichev et al. (2022) offered a model to

optimize the treatment or intensive care of seriously ill Covid-19 patients with plasma proteomics [34]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

By referring to the elaboration of the most recent previous related work by some researchers, the research carried out in this article has novelties that previous researchers have not studied. The novelty of the study lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

III. METHODOLOGY

This study applies two kinds of data mining methods or machine learning algorithms: RF and SVM. The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The development of the application program in this study uses the Python programming language.

The Data Mining Process for Covid-19 patient big data in this study uses CRISP-DM (Cross-Industry Standard Process for Data Mining) process; CRISP-DM is standard data mining. The CRISP-DM process consists of a six-stage [35]; see Figure 1 [36]. Figure 2 shows the data mining process in this study. This research uses confusion matrix and k-fold cross validation in measuring the classification performance of RF and SVM methods.

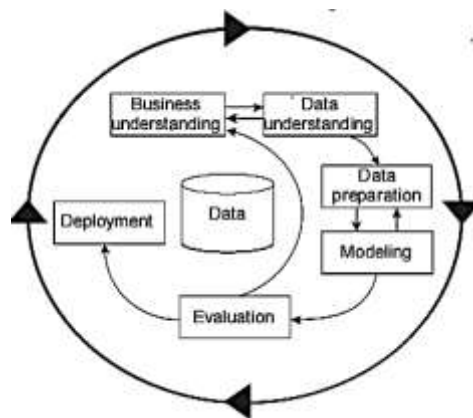


Figure 1. The CRISP-DM Process

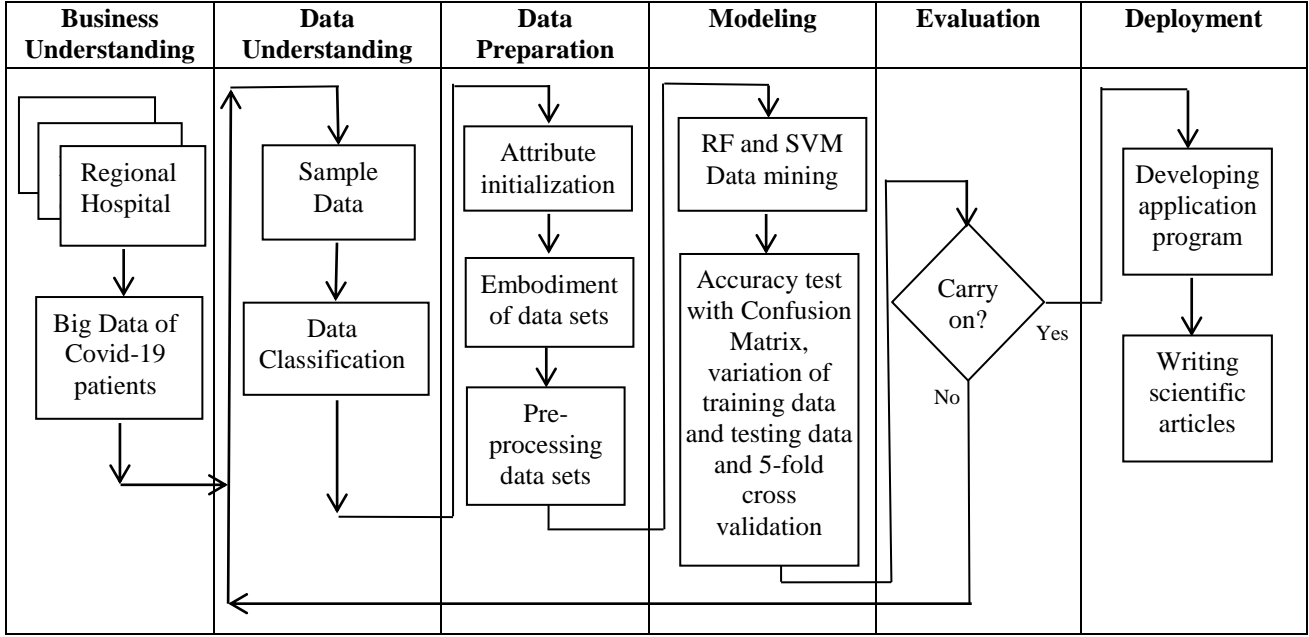


Figure 2. Data Mining Process of Covid-19 Patient Big Data

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected in the form of medical document data from all Covid-19 patients registered at the hospital includes the patient's name, symptoms of the disease, and the treatment status specified. The big data of Covid-19 patients obtained was 117. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each attribute of the treatment status class. The existing attributes and categories are representations of disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 1 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE I. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from other patients, and some patients have similar diagnoses to others. The number of signs and symptoms of the study or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 2).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or G_i where $i = 1, 2, 3 \dots 12$) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 3. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or

symptoms.

TABLE II. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE III. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	yes	no	no	No	No	No	Inpatient
2	yes	yes	no	Yes	No	No	Inpatient
3	yes	no	yes	No	No	No	Inpatient
4	yes	no	no	No	No	No	Inpatient
5	yes	no	no	No	No	No	Inpatient
6	yes	no	no	No	Yes	Yes	Inpatient
7	no	no	yes	No	No	No	Inpatient
..
..
114	no	no	no	No	No	Yes	Self-isolation
115	yes	no	no	No	Yes	No	Inpatient
116	yes	no	no	No	Yes	Yes	Inpatient
117	no	no	no	No	No	No	Inpatient

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1.

In addition, preprocessing the data set is also carried out on the class attribute, namely changing the class attribute with y and changing the value of the self-isolation class attribute with the number 1 and the *inpatient class attribute with the number -1, as shown in Table 4.

TABLE IV. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
..
..
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [37], which has their respective advantages in building application programs [38][39]. The application program built in this research uses the Python programming language to facilitate patient care status prediction.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

The formula of the SVM data mining method is: to form a linear equation from the training data; find the value of w and b, and the value of the classification decision is as follows.

$$S = ((x1, y1), \dots, (xl, yl)) \quad (1)$$

$$yi ((w \cdot xi) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum j^2 \quad (4)$$

$$Entropy(S) = \sum i - pi \times \log_2(pi) \quad (5)$$

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

The format of the confusion matrix table is as shown in Table 5.

TABLE V. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-Validation

K-fold cross-validation was also used to measure the performance of the classification method. K-fold cross-validation is one of the cross-validation tests to assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the K k-fold value. In performance testing of this study with k-fold cross validation, the dataset is partitioned into 5 subsets (k = 5) and allows each subset to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the

training set (Dtrain). In the second fold, the second subset is the validate set the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The performance evaluation of the model carried out is on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is as shown in Table 6. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE VI. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TEST DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% data 90% testing and training data and 10% testing data are as shown in Table 7.

TABLE VII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data		Accuracy		Precision		Recall	
(in %)		(in %)		(in %)		(in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Performance in predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than using the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

Testing the performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models are presented in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

4.6 Deployment

One of the deployments in this research is in the form of making scientific articles on machine learning system models that are produced to be published in reputable scientific articles. Another form of deployment is in the form of making reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 92%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Suggestion for further research is to produce a machine learning system model to predict the death and recovery status of each Covid-19 patient. Another suggestion is to conduct further research using other predictive data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first author, second, and so on as the fifth author.

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Similarity Check _ Covid-19 Patient Care Status Using Random Forest and Support Vector Machine Methods

By Anthony Anggrawan

Covid-19 Patient Care Status Using Random Forest and Support Vector Machine Methods

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Abstract—Covid-19 disease spread throughout the world and caused death globally. Accordingly researchers use many research methods to control the Covid-19 pandemic, one of which is getting more attention is the use of prediction methods. The problem in medical action is not easy to determine the treatment status. So it is not surprising that previous researchers emphasized that the most frequent mistakes were due to inaccuracies in medical decisions. Therefore, this study proposes a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analytics for better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. So, in essence, it is not easy to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of the field of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. In fact, machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [19][20][21]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [22]. For this reason, or that is why this study proposes a machine learning system model for decision-making solutions (predictions) of the treatment status of Covid-19 patients, both inpatient and self-isolation using data mining methods.

Two things machine learning can predict, are classification to predict class membership and regression to show numerical values [12]. While data mining is actually part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in the field of prediction [23]. The

embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting the new data with high accuracy [22].

There are many Data Mining methods, including K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out in order to be able to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [24] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [25] [26], as well as SVM algorithm has a widely known method used for classification [26].

However, it is essential to know the level of accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study, whether the patient should be hospitalized or self-isolated. Therefore, this study also conducted a further testing of the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. Testing the accuracy of predicting the treatment status of Covid-19 patients is carried out on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some of the related works from the latest scientific articles compared with the work in this research article conducted.

- Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [24]. This previous research is different from the research in the article on the research objectives and the object

under study.

- Celestine Iwendi et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [27]. The difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM methods. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.
- Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [28]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, research in this article predicts the treatment status of Covid-19 patients.
- Based on patient clinical data, Boran Hao et al. (2020) developed a model to predict the severity of pneumonia and the level of care required by Covid-19 patients using statistical methods [29]. This previous research differs in the research purpose and method compared to the research in this article.
- Anthony Anggrawan et al. (2021) implemented machine learning to diagnose drug users and types of drug users using Forward Chaining and Certainty Factor methods [22]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.
- Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future by using the Poisson distribution and the gamma distribution [30]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article.
- Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. The research in this article is different from previous research; the difference lies in the research objectives and the research methods used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. The research in this article is not using the DT method but uses the RF and SVM methods.
- Satrio M. Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [31]. This previous study differed in its research objectives from the research in this article. The

previous research predicts patients suffering from Covid-19 or not using the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods.

- Ankit Mehrotraa and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [32]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from the research in this article because this article is an experimental study, not a literature review research.
- Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods used compared to the research in this article. The difference lies in the fact that previous study examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].
- Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [33]. However, the previous research differs in the purpose and the research method compared to the research in this article.
- Vadim Demichev et al. (2022) offered a model to

optimize the treatment or intensive care of seriously Covid-19 patients with plasma proteomics [34]. This previous research is different from the research in the article on the research objectives, research methods, and the object under study.

By referring to the elaboration of the most recent previous related work by some researchers, the research carried out in this article has novelties that previous researchers have not studied. The novelty of the study lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

III. METHODOLOGY

This study applies two kinds of data mining methods or machine learning algorithms: RF and SVM. The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The development of the application program in this study uses the Python programming language.

The Data Mining Process for Covid-19 patient big data in this study uses CRISP-DM (Cross-Industry Standard Process for Data Mining) process; CRISP-DM is standard data mining. The CRISP-DM process consists of a six-stage [35]; see Figure 1 [36]. Figure 2 shows the data mining process in this study. This research uses confusion matrix and k-fold cross validation in measuring the classification performance of RF and SVM methods.



Figure 1. The CRISP-DM Process

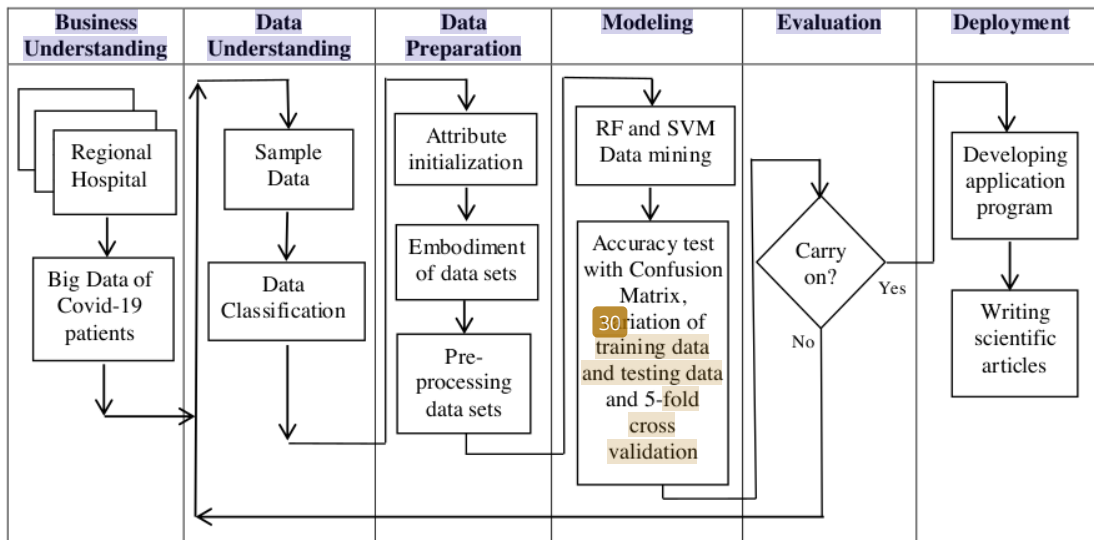


Figure 2. Data Mining Process of Covid-19 Patient Big Data

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected in the form of medical document data from all Covid-19 patients registered at the hospital includes the patient's name, symptoms of the disease, and the treatment status specified. The big data of Covid-19 patients obtained was 117. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each attribute of the treatment status class. The existing attributes and categories are representations of disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 1 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE I. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from other patients and some patients have similar diagnoses to others. The number of signs and symptoms of the study or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 2).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or Gi where i = 1, 2, 3 ...12) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 3. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or

symptoms.

TABLE II. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE III. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	yes	no	no	No	No	No	Inpatient
2	yes	yes	no	Yes	No	No	Inpatient
3	yes	no	yes	No	No	No	Inpatient
4	yes	no	no	No	No	No	Inpatient
5	yes	no	no	No	No	No	Inpatient
6	yes	no	no	No	Yes	Yes	Inpatient
7	no	no	yes	No	No	No	Inpatient
..
..
114	no	no	no	No	No	Yes	Self-isolation
115	yes	no	no	No	Yes	No	Inpatient
116	yes	no	no	No	Yes	Yes	Inpatient
117	no	no	no	No	No	No	Inpatient

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1.

In addition, preprocessing the data set is also carried out on the class attribute, namely changing the class attribute with y and changing the value of the self-isolation class attribute with the number 1 and the *inpatient class attribute with the number -1, as shown in Table 4.

TABLE IV. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
..
..
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

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The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [37], which has their respective advantages in building application programs [38][39]. The application program built in this research uses the Python programming language to facilitate patient care status prediction.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

The formula of the SVM data mining method is: to form a linear equation from the training data; find the value of w and b, and the value of the classification decision is as follows.

$$S = ((x1, y1), \dots, (xl, yl)) \quad (1)$$

$$yi ((w \cdot xi) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum p_i^2 \quad (4)$$

$$Entropy(S) = \sum p_i - p_i \times \log_2(p_i) \quad (5)$$

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

The format of the confusion matrix table is as shown in Table 5.

TABLE V. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-Validation

K-fold cross-validation was also used to measure the performance of the classification method. K-fold cross-validation is one of the cross-validation tests to assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the K value. In performance testing of this study with k-fold cross validation, the dataset is partitioned into 5 subsets (k = 5) and allows each subset to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken obviously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the

training set (Dtrain). In the second fold, the second subset is the validate set the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The performance evaluation of the model carried out is on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is as shown in table 6. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE VI. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TEST DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% testing data and 90% training data and 10% testing data are as shown in Table 7.

TABLE VII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)	Accuracy (in %)	Precision (in %)	Recall (in %)	
Training Testing	RF SVM	RF SVM	RF SVM	
50	50	95 97	97 96	91 96
60	40	96 91	97 90	93 90
70	30	94 92	96 91	92 90
80	20	96 92	94 89	97 94
90	10	92 83	75 67	95 91
Average		95 91	92 87	94 92

Performance in predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than using the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

Testing the performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models are presented in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

4.6 Deployment

One of the deployments in this research is in the form of making scientific articles on machine learning system models that are produced to be published in reputable scientific articles. Another form of deployment is in the form of making reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 92%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Suggestion for further research is to produce a machine learning system model to predict the death and recovery status of each Covid-19 patient. Another suggestion is to conduct further research using other predictive data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first author, second, and so on as the fifth author.

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By Anthony Anggrawan

Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

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Abstract—Covid-19 has become a global pandemic that causes many deaths, so medical treatment for Covid-19 patients gets special attention, whether hospitalized or self-isolated. However, the problem in medical action is not easy, and the most frequent mistakes are due to inaccuracies in medical decision-making. Meanwhile, machine learning predict with high accuracy. For that, or that's why this study aims to propose a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients. The implication is that RF machine learning can help/replace the role of medical experts in predicting the patient's care status.

Index Terms—data mining, random forest, support vector machine, prediction, Covid-19, machine learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1]-[3]. It is the cause of the global health crisis [4]-[6], which is not because of its high-speed transmission [5], [7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analytics for better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for

Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these hospitalized Covid-19 patients are receiving intensive medical care from doctors.

In essence, the care status for Covid-19 patients is self-isolation for patients with non-severe illness status and hospitalization for patients who are seriously ill and at critical risk or cause death. Hospitals or medical doctors take various ways to reduce the number of deaths of Covid-19 patients, including by regulating the status of services in hospitals. Expert doctors recommend that patients self-isolate or should be hospitalized after analyzing the patient's medical data. Determining the level of care for Covid-19 patients is a form of medical treatment or treatment for Covid-19 patients to get proper treatment or care.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both in hospital and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16], [17]; moreover, with advances in information and communication technology [18], [19], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [20]-[22]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more

accurately [13], and provide high-accuracy prediction results [23]. For this reason (or why), this study's objective is to propose a machine learning system model for decision-making solutions (predictions) for the treatment status of Covid-19 patients, both inpatient and self-isolation, using data mining methods.

The implication is that the proposed machine learning system model can help and even replace the role of medical experts (specialist doctors) in making medical decisions for Covid-19 patients, whether hospitalization or self-isolation. Machine learning performs tasks like a medical specialist in determining the results of the diagnosis of the infection status of Covid-19 patients based on medical data of Covid-19 patients. Furthermore, machine learning can work tirelessly, time and place, and has intelligence like an expert, so it is not surprising that previous research confirms that intelligent machines can make superior decisions to experts because humans have a human error factor [24].

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [25]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting new data with high accuracy [23]. After all, predicting individuals with symptoms of being infected with Covid-19 mandates machine learning (application-based) and contributes to effectively isolating Covid-19 patients [26].

Big data demands large storage media [27]. However, big data is no longer traditionally processed [28]. Instead, today's big data processing relies on machines that can provide systematic results [29]. Big data storage is generally on a computer server with a large storage capacity. Still, some also make it happen by renting online cloud data storage services such as Amazon Simple Storage Service (Amazon S3) and Google Cloud [30]. Cloud facilitates cost-effective big data storage and analysis [30].

Big data processing techniques in data mining include several stages: target data, preprocessed data, data mining, and evaluation/analysis of mining results [30]. Target data and preprocessed data are the processes of extracting raw data from big data [30]. Target data is to select the required data (sample data) and classify data. The preprocessed data is to prepare data sets for data mining, including cleaning up incomplete data, duplicate data, and converting string data into numeric coded data. Finally, data mining and evaluation results extract hidden information by applying data mining methods suitable for the objectives and analyzing them. [88]

Many Data Mining methods include K-Means, Naive Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There

are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [31] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [32], [33], as well as SVM algorithm has a widely known technique used for classification [33]. It is why this study uses SVM and RF to classify treatment status. Given that the SVM and RF machine learning algorithms are both popularly used by many researchers, the SVM and RF machine learning algorithms are the most appropriate combination used in research, including research to classify and predict the treatment status of Covid-19 patients.

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods. [5]

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some related works from the latest scientific articles compared with the work in this research article.

Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [31]. This previous research is different from the research in the article on the research objectives and the object under study. In the meantime, Ceine Iwendi *et al.* (2020) proposed the Random Forest model to predict disease severity of Covid-19 patients [34]. The difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this

article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.

Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [35]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, the research in this article predicts the treatment status of Covid-19 patients.

Based on patient clinical data, using statistical methods, Boran Hao *et al.* (2020) developed a model to predict pneumonia severity in Covid-19 patients using the Natural Language Processing tool [36]. However, this previous research differs in the research purpose and way compared to the research in this article. Meanwhile, Anthony Anggrawan *et al.* (2021) implemented machine learning to diagnose drug users and types of drug-using forward Chaining and Certainty Factor methods [23]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.

Hongwei Zhao *et al.* (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson and Gamma distribution [37]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article. In the meantime, Bassam Mahboub *et al.* (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research objectives and techniques used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. This article's research does not use the DT method but uses the RF and SVM methods.

Soham Guhathakurata *et al.* (2021) predicted whether a person is infected with Covid-19 or not using SVM [38]. However, this previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not utilizing the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods. At the same time, Ankit Mehrotra and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [39]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However,

it is different from this article's research because it is an experimental study, not a literature review.

Pratiyush Guleria *et al.* (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods compared to this article's research. The difference is that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].

Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [40]. However, the previous research differs in the purpose and way compared to this article's research. In contrast, Vadim Demichev *et al.* (2022) offered a model to optimize the treatment or intensive care of seriously ill Covid-19 patients with plasma proteomics [41]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

Table I compares some of the most recent previous related work with the work carried out in this study. By referring to the elaboration of the most recent last related work by some researchers, the research carried out in this article has novelties (from the prior research gap) that previous researchers have not studied. In essence, the gap in earlier research is that no one has researched machine learning models to predict the inpatient status or self-isolation of Covid-19 patients by involving RF and SVM algorithms. In addition, the 12 criteria used to indicate the treatment status of Covid-19 patients are entirely different from previous similar studies (as shown in Table I in the Criteria/Attributes column). So, this study's originality lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

III. METHODOLOGY

This study applies two data mining methods or machine learning algorithms: RF and SVM. The big data is on Covid-19 patients from a regional hospital in Mataram, Indonesia. The significant data source used in this study is primary data from patient medical records/documents. The attributes of the patient's disease symptoms and care status classes amount to thousands of patient medical record data. Patient datasets containing non-Covid-19 and duplicate and incomplete Covid-19

patient data are removed, so only data is left as a dataset for data mining processes. Medical records data of disease symptoms obtained from string data is then converted into numeric data. The development of the

application program in this study uses the Python computer programming language.

TABLE I. COMPARISON OF THIS ARTICLE'S WORK WITH SOME PREVIOUS RELATED WORKS

Research by	Research methods			Criteria/Attributes		Research Object	Accuracy Test
	RF	SVM	ML	Number	Name		
Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) [31]	Yes	Yes	Yes	10	58 Coastal Aerosol, Blue, Green, Red, Vegetation Red Edge, NIR, Narrow NIR, Water vapor, SWIR-Cirrus, SWIR	Satellite remote sensing images	Yes
Celestine Iwendi <i>et al.</i> (2020) [34]	Yes	No	Yes	6	65 Symptom1, symptom2, symptom3, symptom4, symptom5, symptom6	Illness severity of Covid-19 patients	No
Chelvian Aroef, Yuda Rivian, and Zuherman Rustam (2020) [35]	Yes	Yes	Yes	9	75 Age, Body Mass Index (BMI), Glucose, Insulin, Homa, Leptin, Adiponectin, Resistin, MCP 1	Breast cancer	Yes
Boran Hao <i>et al.</i> (2020) [36]	No	No	No	10	75 Radiology Opacities, Respiratory Rate, Age, Fever Male, Albumin, Anion Gap, SpO2, LDH, Calcium	The severity of pneumonia in Covid-19 patients	Yes
Anthony Anggrawan <i>et al.</i> (2021) [23]	No	No	Yes	27	116 Out of breath, Anxious, Nausea, Diarrhea, Convulsions, Easily angry, Depression, Sleep patterns change, Sweating, Chills, Shaking, Insomnia, Fast heart rate, Blood pressure rises, Difficult to focus, Difficult to rest, Weight loss, Dry mouth, Blurred vision, Changed skin color, Constipation, Stomachache, Drowsiness, Itching, Difficulty urinating, Mood swings, Dizziness	Drug users and types of drug-using	Yes
Hongwei Zhao <i>et al.</i> (2021) [37]	No	No	No	0	-	Number of cases of Covid-19 patients	No
Bassam Mahboub <i>et al.</i> (2021) [8]	No	No	Yes	5	31 Age, Gender, Nationality, Blood group, BMI	The treatment period for Covid-19 patient	Yes
Soham Guhathakurata <i>et al.</i> (2021) [38]	No	Yes	Yes	8	Temp, Breathing rate, Hypertension, Heartbeat rate (HBR), Acute respiratory disease syndrome (ARDS), Chest pain, Heart disease, Cough with sputum (CWS)	Predicting whether patients are infected with Covid-19 or not	No
Ankit Mehrotraa and Reeti Agarwal (2021) [39]	No	No	No	0	-	Discussing the data mining method's role in the Covid-19 pandemic	No
Pratiyush Guleria <i>et al.</i> (2022) [14]	No	Yes	Yes	0	-	The death rate of Covid-19 patient	Yes
Anthony Anggrawan, <i>et al.</i> (2022) [40]	No	No	Yes	6	2 Achievement indicators, achievement points, recommendation, organizational activity, semester level, and completeness of documents	Scholarship recipient	Yes
Vadim Demichev <i>et al.</i> (2022) [41]	No	No	No	0	-	Optimization of treatment for Covid-19 patients	Yes
Our research	Yes	Yes	Yes	12	Pneumonia, ARDS, CHF, AKI, CAD, Dyspnea, NSTEMI, ADHF, HHD, Febris, Anosmia, Ageusia	Care status of Covid-19 patients, whether inpatient or self-isolation	Yes

Note: ML = Machine Learning

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods. The data mining process in this research uses CRISP-DM (Cross-Industry Standard Process for Data Mining). CRISP-DM is a standard data mining process. The process in CRISP-DM comprises a six-stage [42], as shown in Fig. 1 [43]. Fig. 2 shows the process carried out at each stage of CRISP-DM.

In Figure 2, business understanding is the stage of sorting out thousands of hospital patient medical data to collect the required patient data. The next stage is understanding the data collected as representative data

for Covid-19 patients. This Covid-19 patient data classifies the patient's signs and symptoms and treatment status, which needs further processing at the next stage. The next stage is the data preparation stage, which essentially determines the attributes of the names of signs and symptoms of Covid-19 patients. The embodiment of the dataset containing knowledge according to treatment status refers to the signs and symptoms that the patient has (marked with Yes) or does not have (marked with No). The next thing to do is preprocess the dataset, changing the category value of the symptom attribute and the class attribute with the number 1 and the number -1. The process of extracting raw data obtained in the

previous process stage is data that is further processed by data mining methods or used as learning machine learning data at the modeling stage. So that machine learning can predict. The next stage is the evaluation stage, namely, knowing the predictive reliability of the data mining or machine learning method. Then the last stage is the deployment stage to disseminate research results so that they are helpful for implementation by various parties, especially hospitals and other professionals, in the form of developing application programs and scientific articles.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

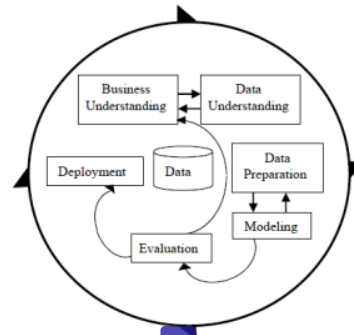


Figure 1. The CRISP-DM process.

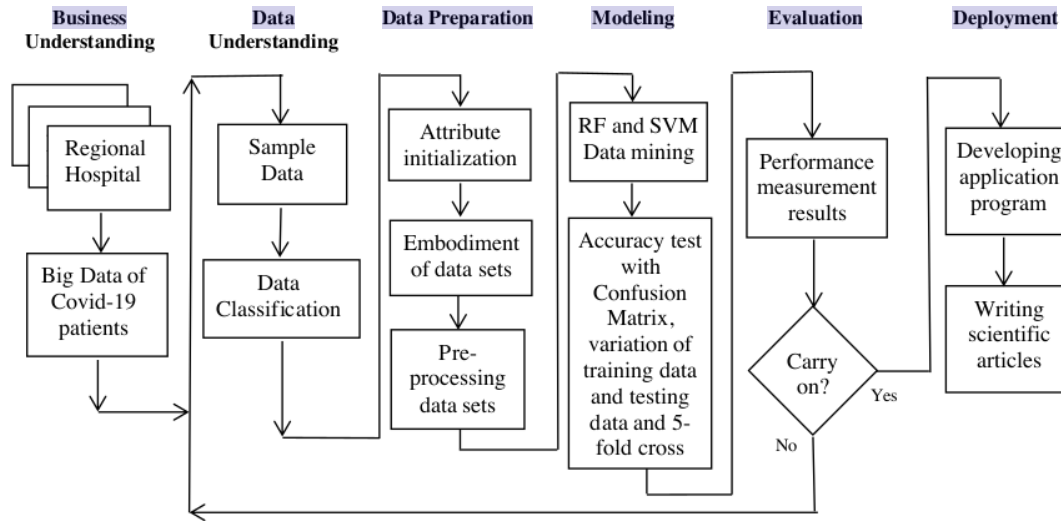


Figure 2. Data mining process of Covid-19 patient big data.

IV. RESULT AND DISCUSSION

A. Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected from medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. There are thousands of data on Covid-19 patients. The patient dataset containing incomplete data and non-covid-19 was omitted or ignored. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria of patient medical data is used in training and testing the prediction model

proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

TABLE II. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

B. Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data

representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table II shows the association between signs and symptoms of disease and treatment status in the study data set.

C. Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses. There were 117 patients with Covid-19 [95] had a different diagnosis from the others. In this study, the number of signs and symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table III).

TABLE III. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or Gi where i = 1, 2,3 ...12) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table IV. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or symptoms.

TABLE IV. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	Yes	No	No	No	No	No	Inpatient
2	Yes	Yes	No	Yes	No	No	Inpatient
3	Yes	No	Yes	No	No	No	Inpatient
4	Yes	No	No	No	No	No	Inpatient
5	Yes	No	No	No	No	No	Inpatient
6	Yes	No	No	No	Yes	Yes	Inpatient
7	No	No	Yes	No	No	No	Inpatient
..
..
114	No	No	No	No	No	Yes	Self-isolation
115	Yes	No	No	No	Yes	No	Inpatient
116	Yes	No	No	No	Yes	Yes	Inpatient
117	No	No	No	No	No	No	Inpatient

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1. In addition, dataset preprocessing is also carried out on class attributes, namely changing the independent isolation

class attribute category with the number 1 and the inpatient class attribute category with the number -1, as shown in Table V.

TABLE V. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
..
..
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

D. Modeling

The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, known various programming language [44], which has their respective advantages in building application programs [45], [46]. The application program built in this research uses the Python programming language to facilitate patient care status prediction (as show in Table VI).

TABLE VI. Use of HYPERPARAMETER on Svm and Rf Method

Classifier Method	Hyperparameter	Value
SVM	C	1
	Kernel	Rbf
	Degree	3
	Gamma	Scale
	Coef	0
	Tol	0.000
RF	74_max_iter	-1
	n_estimators	100
	Criterion	Gini
	Max_depth	None
	Min_samples_split	2
Min_samples_leaf	1	

1) SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: 1) Forming a linear equation from the training data that has gone through the preprocessing stage; 2) Finding the values of w and b by means of elimination and substitution of linear equations; 3) Finding the value of the classification decision with the function.

In SVM, there are two implementation models: mathematical programming techniques and kernel functions. This study applies kernel functions and focuses on classifying two categories of class attributes. The class attribute is a treatment for $y_i = +1, -1$. The formula of the SVM data mining method is: 1) to form a linear equation

from the training data; 2) find the value of w and b , and 3) the value of the classification decision is as follows.

$$S = \{(x_1, y_1), \dots, (x_l, y_l)\} \quad (1)$$

$$y_i ((w \cdot x_i) + b) \geq 1, i = 1 \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S = set; x = attribute; y = class; w = weight; b = bias

2) RF data mining method

The process of realizing the classification using the RF data mining method is as follows: 1) Generating a random subset of data; 2) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and 3) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

RF uses bootstrap samples from training data to create a tree from a randomly selected subset. The chosen predictor is a candidate for splitting the decision tree. The results of the category predictions from the treatment class based on the results of the highest voting were chosen as the final prediction results. The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_{i=1}^c p_i^2 \quad (4)$$

$$Entropy(S) = \sum_{i=1}^c -p_i \times \log_2(p_i) \quad (5)$$

Description:

S = Set of cases

p_i = the proportion of case i to the Set of cases

3) Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected as positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected as negative. Meanwhile, the calculation of accuracy, prediction, and Recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

Accuracy states the closeness of the measurement results to the actual value, while Precision shows how close the difference in the measurement results is on repeated measurements. On the other hand, Recall states the level of success in retrieving information. Precision and Recall are

necessary because Precision denotes a measure of quality, and Recall denotes a measure of quantity.

Measurement of accuracy is based on the ratio between the correct predictions (positive and negative) with the overall data. In contrast, precision measurements are based on the percentage of true positive predictions compared to overall positive predicted outcome. Meanwhile, the recall measurement is based on the ratio of true positive predictions compared to the general actual positive data.

The format of the confusion matrix table is as shown in Table VII. The results of the predictions of the SVM and RF methods are shown in Tables VIII and IX.

TABLE VII. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

TABLE VIII. CONFUSION MATRIX OF SVM

		Prediction	
		Self-isolation	Inpatient
Actual	Class		
	Self-isolation	4	0
	Inpatient	2	12

TABLE IX. CONFUSION MATRIX OF RF

		Prediction	
		Self-isolation	Inpatient
Actual	Class		
	Self-isolation	4	0
	Inpatient	1	13

4) K-fold Cross-validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation helps assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each subgroup to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the training set (Dtrain). In the second fold, the second subset is the validate set, the remaining subset is the training set, and so on until the 5th fold.

E. Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

1) Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in

90]le X and Figure 3. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE X. . SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TESTING DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

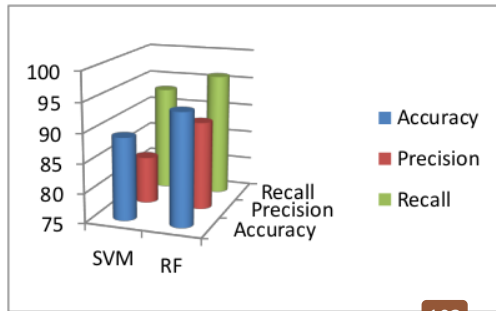


Figure 3. System model performance testing with 85% of training data and 15% of testing data.

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% data 90% testing and training data and 10% testing data are as shown in Table XI.

TABLE XI. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)		Accuracy (in %)		Precision (in %)		Recall (in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix.

2) Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table XII and Fig. 4.

TABLE XII. PREDICTION PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

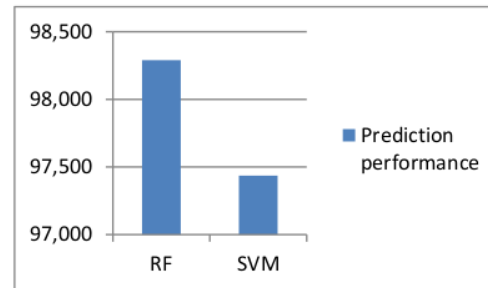


Figure 4. Predictive performance testing using K-fold cross-validation.

F. Deployment

One of the deployments in this research is making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Thus, the results obtained can be developed and become the knowledge of many parties as a responsibility for the correctness of the effects of research carried out as professional researchers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 90%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%. The research result implication is that RF machine learning can help or replace the role of medical personnel in predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, with high accuracy.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Further research needs to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion for future study is: to conduct further research using other data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients and various other diseases, to build a system that not only predicts but also performs clustering, association, and estimates of

various other fields of science including patients' care status, with a combination of machine learning and the Internet of Things.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first, second, and so on as the fifth author.

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By Anthony Anggrawan

Covid-19 Patient Care Status Using Random Forest and Support Vector Machine Methods

Anthony Anggrawan, Mayadi, Christofer Satria, Bambang Krismono Triwijoyo and Ria Rismayati
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Abstract—Covid-19 disease spread throughout the world and caused death globally. Accordingly researchers use many research methods to control the Covid-19 pandemic, one of which is getting more attention is the use of prediction methods. The problem in medical action is not easy to determine the treatment status. So it is not surprising that previous researchers emphasized that the most frequent mistakes were due to inaccuracies in medical decisions. Therefore, this study proposes a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analytics for better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. So, in essence, it is not easy to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of the field of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. In fact, machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [19][20][21]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [22]. For this reason, or that is why this study proposes a machine learning system model for decision-making solutions (predictions) of the treatment status of Covid-19 patients, both inpatient and self-isolation using data mining methods.

Two things machine learning can predict, are classification to predict class membership and regression to show numerical values [12]. While data mining is actually part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in the field of prediction [23]. The

embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting the new data with high accuracy [22].

There are many Data Mining methods, including K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out in order to be able to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [24] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [25] [26], as well as SVM algorithm has a widely known method used for classification [26].

However, it is essential to know the level of accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study, whether the patient should be hospitalized or self-isolated. Therefore, this study also conducted a further [62] of the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. Testing the accuracy of predicting the treatment status of Covid-19 patients is carried out on both RF and SVM machine learning methods. 4

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some of the related works from the latest scientific articles compared with the work in this research article conducted.

- Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [24]. This previous research is different from the research in the article on the research objectives and the object

under study.

- Celestia Iwendu et al. (2020) proposed the Random Forest model to predict disease severity of Covid-19 patients [27]. The difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM methods. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.
- Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [28]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, research in this article predicts the treatment status of Covid-19 patients. 19
- Based on patient clinical data, Boran Hao et al. (2020) developed a model to predict the severity of pneumonia and the level of care required by Covid-19 patients using statistical methods [29]. This previous research differs in the research purpose and method compared to the research in this article.
- Anthony Anggrawan et al. (2021) implemented machine learning to diagnose drug users and types of drug users using Forward Chaining and Certainty Factor methods [22]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM. 55
- Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future by using the Poisson distribution and the gamma distribution [30]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article. 19
- Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. The research in this article is different from previous research; the difference lies in the research objectives and the research methods used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. The research in this article is not using the DT method but uses the RF and SVM methods.
- Srim Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [31]. This previous study differed in its research objectives from the research in this article. The

previous research predicts patients suffering from Covid-19 or not using the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods.

- Ankit Mehrotraa and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [32]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from the research in this article because this article is an experimental study, not a literature review research.
- Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods used compared to the research in this article. The difference lies in the fact that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].
- Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [33]. However, the previous research differs in the purpose and the research method compared to the research in this article.
- Vadim Demichev et al. (2022) offered a model to

optimize the treatment or intensive care of seriously Covid-19 patients with plasma proteomics [34]. This previous research is different from the research in the article on the research objectives, research methods, and the object under study.

By referring to the elaboration of the most recent previous related work by some researchers, the research carried out in this article has novelties that previous researchers have not studied. The novelty of the study lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

III. METHODOLOGY

This study applies two kinds of data mining methods or machine learning algorithms: RF and SVM. The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The development of the application program in this study uses the Python programming language.

The Data Mining Process for Covid-19 patient big data in this study uses CRISP-DM (Cross-Industry Standard Process for Data Mining) process; CRISP-DM is standard data mining. The CRISP-DM process consists of a six-stage process [35]; see Figure 1 [36]. Figure 2 shows the data mining process in this study. This research uses confusion matrix and k-fold cross validation in measuring the classification performance of RF and SVM methods.

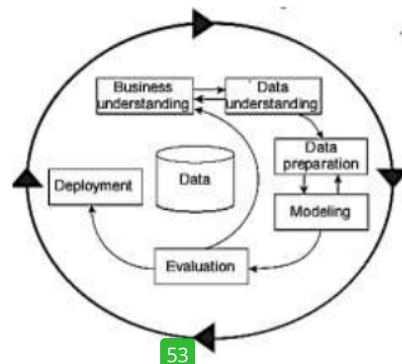


Figure 1. The CRISP-DM Process

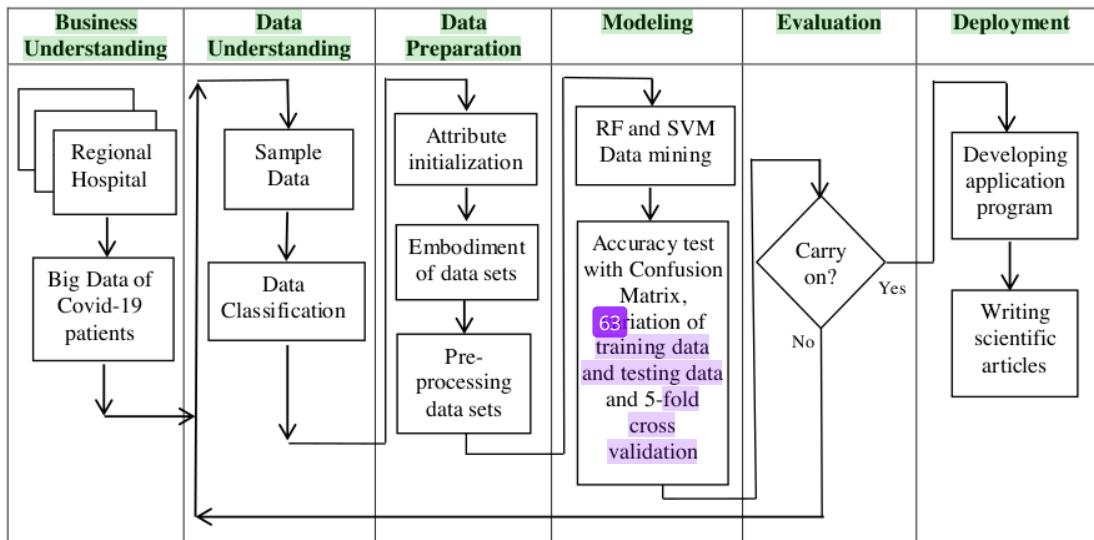


Figure 2. Data Mining Process of Covid-19 Patient Big Data

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected in the form of medical document data from all Covid-19 patients registered at the hospital includes the patient's name, symptoms of the disease, and the treatment status specified. The big data of Covid-19 patients obtained was 117. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each attribute of the treatment status class. The existing attributes and categories are representations of disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 1 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE I. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from other patients and some patients have similar diagnoses to others. The number of signs and symptoms of the study or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 2).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or Gi where i = 1, 2, 3 ...12) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 3. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or

symptoms.

TABLE II. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE III. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	yes	no	no	No	No	No	Inpatient
2	yes	yes	no	Yes	No	No	Inpatient
3	yes	no	yes	No	No	No	Inpatient
4	yes	no	no	No	No	No	Inpatient
5	yes	no	no	No	No	No	Inpatient
6	yes	no	no	No	Yes	Yes	Inpatient
7	no	no	yes	No	No	No	Inpatient
..
..
114	no	no	no	No	No	Yes	Self-isolation
115	yes	no	no	No	Yes	No	Inpatient
116	yes	no	no	No	Yes	Yes	Inpatient
117	no	no	no	No	No	No	Inpatient

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1.

In addition, preprocessing the data set is also carried out on the class attribute, namely changing the class attribute with y and changing the value of the self-isolation class attribute with the number 1 and the *inpatient class attribute with the number -1, as shown in Table 4.

TABLE IV. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
..
..
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

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The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [37], which has their respective advantages in building application programs [38][39]. The application program built in this research uses the Python programming language to facilitate patient care status prediction.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

The formula of the SVM data mining method is: to form a linear equation from the training data; find the value of w and b, and the value of the classification decision is as follows.

$$S = ((x1, y1), \dots, (xl, yl)) \quad (1)$$

$$yi ((w \cdot xi) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum^2 \quad (4)$$

$$Entropy(S) = \sum i - pi \times \log_2(pi) \quad (5)$$

4.4.3 Confusion matrix

This research uses a confusion matrix to measure performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

The format of the confusion matrix table is as shown in Table 5.

TABLE V. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-Validation

K-fold cross-validation was also used to measure the performance of the classification method. K-fold cross-validation is one of the cross-validation tests to assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the K value. In performance testing of this study with k-fold cross validation, the dataset is partitioned into 5 subsets (k = 5) and allows each subset to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken obviously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the

training set (Dtrain). In the second fold, the second subset is the validate set the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The performance evaluation of the model carried out is on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with

Confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is as shown in table 6. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE VI. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TEST DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% testing data and 90% training data and 10% testing data are as shown in Table 7.

TABLE VII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)	Accuracy (in %)	Precision (in %)	Recall (in %)	
Training Testing	RF SVM	RF SVM	RF SVM	
50	50	95 97	97 96	91 96
60	40	96 91	97 90	93 90
70	30	94 92	96 91	92 90
80	20	96 92	94 89	97 94
90	10	92 83	75 67	95 91
Average		95 91	92 87	94 92

Performance in predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than using the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

Testing the performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models are presented in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

4.6 Deployment

One of the deployments in this research is in the form of making scientific articles on machine learning system models that are produced to be published in reputable scientific articles. Another form of deployment is in the form of making reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 92%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Suggestion for further research is to produce a machine learning system model to predict the death and recovery status of each Covid-19 patient. Another suggestion is to conduct further research using other predictive data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first author, second, and so on as the fifth author.

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Machine Learning to Predict the Treatment Status of Covid-19 Patients

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Abstract—Covid-19 disease spread throughout the world and caused death globally. Accordingly, researchers use many research methods to control the Covid-19 pandemic, one of which is getting more attention is the use of prediction methods. However, the problem in medical action is not easy to determine the treatment status. So it is not surprising that previous researchers emphasized that the most frequent mistakes were due to inaccuracies in medical decisions. Therefore, this study proposes a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not only because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analytics for better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these hospitalized Covid-19 patients are receiving intensive medical care from doctors.

Errors in decision-making often occur because

decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [19][20][21]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [22]. For this reason, or that is why this study proposes a machine learning system model for decision-making solutions (predictions) of the treatment status of Covid-19 patients, both inpatient and self-isolation using data mining methods.

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [23]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting the new data with high accuracy [22].

Many Data Mining methods include K-Means,

Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out in order to be able to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [24] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [25] [26], as well as SVM algorithm has a widely known technique used for classification [26].

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some of the related works from the latest scientific articles compared with the work in this research article.

- Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [24]. This previous research is different from the research in the article on the research objectives and the object under study.
- Celestine Iwendi et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [27]. The difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.
- Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [28]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, the research in this article predicts the treatment status of Covid-19 patients.
- Based on patient clinical data, Boran Hao et al. (2020) developed a model to predict pneumonia severity and the level of care required by Covid-19 patients using statistical methods.[29]. This previous research differs in the research purpose and way compared to the research in this article.
- Anthony Anggrawan et al. (2021) implemented machine learning to diagnose drug users and types of drug-using Forward Chaining and Certainty Factor methods [22]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.
- Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson distribution and the gamma distribution [30]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article.
- Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research objectives and methods used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. This article's research does not use the DT method but the RF and SVM methods.
- Soham Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [31]. This previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not using the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods.
- Ankit Mehrotraa and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [32]. This previous research is a literature review study that concludes that the Data

Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from the research in this article because this article is an experimental study, not a literature review research.

- Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods used compared to the research in this article. The difference lies in the fact that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].
- Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [33]. However, the previous research differs in the purpose and method compared to this article's research.
- Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of seriously ill Covid-19 patients with plasma proteomics [34]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

By referring to the elaboration of the most recent

previous related work by some researchers, the research carried out in this article has novelties that previous researchers have not studied. The novelty of the study lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

III. METHODOLOGY

This study applies two kinds of data mining methods or machine learning algorithms: RF and SVM. The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The development of the application program in this study uses the Python programming language.

The Data Mining Process for Covid-19 patient big data in this study uses CRISP-DM (Cross-Industry Standard Process for Data Mining) process; CRISP-DM is standard data mining. The CRISP-DM process comprises a six-stage [35]; see Figure 1 [36]. Figure 2 shows the data mining process in this study. This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

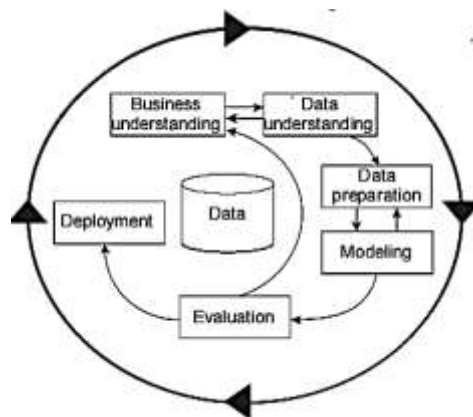


Figure 1. The CRISP-DM Process

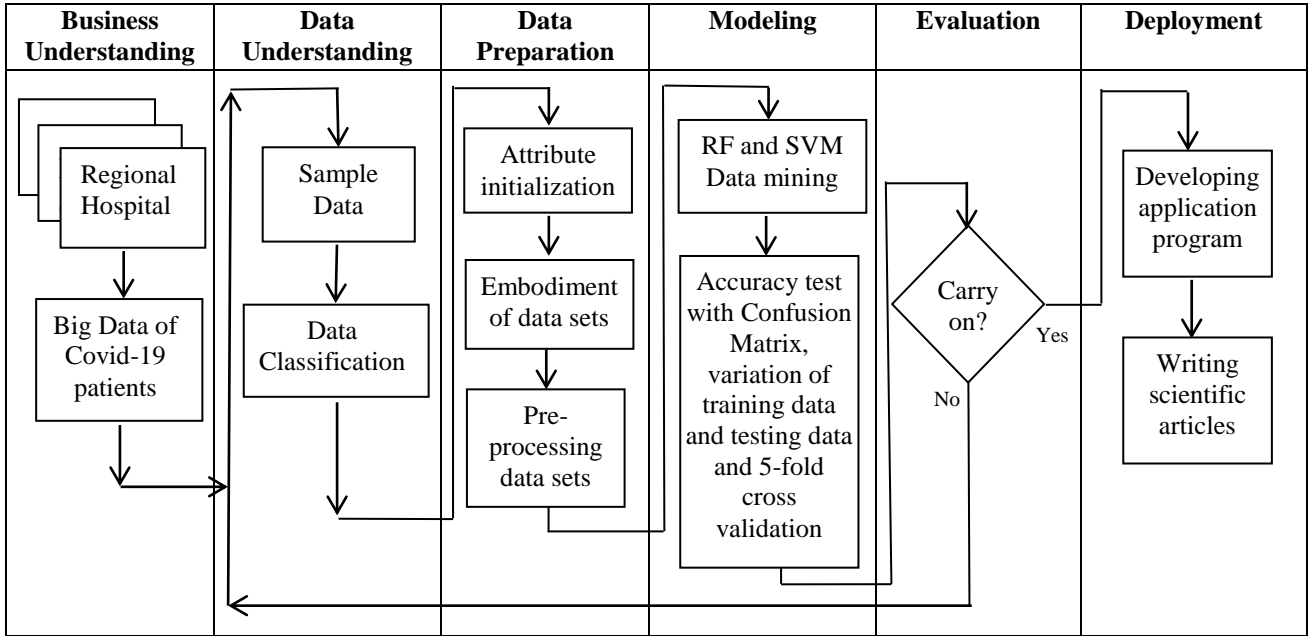


Figure 2. Data Mining Process of Covid-19 Patient Big Data

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected in the form of medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. The big data of Covid-19 patients obtained was 117. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 1 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE I. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses to others. In this study, the number of signs and symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 2).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or G_i where $i = 1, 2, 3 \dots 12$) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 3. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or

symptoms.

TABLE II. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE III. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	yes	no	no	No	No	No	Inpatient
2	yes	yes	no	Yes	No	No	Inpatient
3	yes	no	yes	No	No	No	Inpatient
4	yes	no	no	No	No	No	Inpatient
5	yes	no	no	No	No	No	Inpatient
6	yes	no	no	No	Yes	Yes	Inpatient
7	no	no	yes	No	No	No	Inpatient
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114	no	no	no	No	No	Yes	Self-isolation
115	yes	no	no	No	Yes	No	Inpatient
116	yes	no	no	No	Yes	Yes	Inpatient
117	no	no	no	No	No	No	Inpatient

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1.

In addition, preprocessing the data set is also carried out on the class attribute, namely changing the class attribute with y and changing the value of the self-isolation class attribute with the number 1 and the *inpatient class attribute with the number -1, as shown in Table 4.

TABLE IV. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
..
..
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [37], which has their respective advantages in building application programs [38][39]. The application program built in this research uses the Python programming language to facilitate patient care status prediction.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

The formula of the SVM data mining method is: to form a linear equation from the training data; find the value of w and b, and the value of the classification decision is as follows.

$$S = ((x1, y1), \dots, (xl, yl)) \quad (1)$$

$$yi ((w \cdot xi) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum j^2 \quad (4)$$

$$Entropy(S) = \sum i - pi \times \log_2(pi) \quad (5)$$

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

The format of the confusion matrix table is as shown in Table 5.

TABLE V. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-Validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation is helpful in assessing the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold value. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each subgroup to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the

training set (Dtrain). In the second fold, the second subset is the validate set the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 6. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE VI. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TEST DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% data 90% testing and training data and 10% testing data are as shown in Table 7.

TABLE VII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data		Accuracy		Precision		Recall	
(in %)		(in %)		(in %)		(in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

4.6 Deployment

One of the deployments in this research is in the form of making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 92%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Suggestions for further research are the need to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion is to conduct further research using other predictive data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first author, second, and so on as the fifth author.

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Similarity Check_Machine Learning Model to Predict the Treatment Status of Covid-19 Patients

By Anthony Anggrawan

Machine Learning to Predict the Treatment Status of Covid-19 Patients

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Abstract—Covid-19 disease spread throughout the world and caused death globally. Accordingly, researchers use many research methods to control the Covid-19 pandemic, one of which is getting more attention is the use of prediction methods. However, the problem in medical action is not easy to determine the treatment status. So it is not surprising that previous researchers emphasized that the most frequent mistakes were due to inaccuracies in medical decisions. Therefore, this study proposes a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analysis or better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these hospitalized Covid-19 patients are receiving intensive medical care from doctors.

Errors in decision-making often occur because

decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [19][20][21]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [22]. For this reason, or that is why this study proposes a machine learning system model for decision-making solutions (predictions) of the treatment status of Covid-19 patients, both inpatient and self-isolation using data mining methods.

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [23]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting the new data with high accuracy [22].

Many Data Mining methods include K-Means,

Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out in order to be able to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [24] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [25] [26], as well as SVM algorithm has a widely known technique used for classification [26].

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some of the related works from the latest scientific articles compared with the work in this research article.

- Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [24]. This previous research is different from the research in the article on the research objectives and the object under study.
- Celestine Iwendi et al. (2020) proposed the Random Forest model to predict disease severity of Covid-19 patients [27]. The difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.
- Chelvia Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [28]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, research in this article predicts the treatment status of Covid-19 patients.
- Based on patient clinical data, Boran Hao et al. (2020) developed a model to predict pneumonia severity and the level of care required by Covid-19 patients using statistical methods.[29]. This previous research differs in the research purpose and way compared to the research in this article.
- Anthony Anggrawan et al (2021) implemented machine learning to diagnose drug users and types of drug-using Ford Chaining and Certainty Factor methods [22]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.
- Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson distribution and the gamma distribution [30]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article.
- Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research objectives and methods used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. This article's research does not use the DT method but the RF and SVM methods.
- Srim Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [31]. This previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not using the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods.
- Ankit Mehrotra and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [32]. This previous research is a literature review study that concludes that the Data

Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from the research in this article because this article is an experimental study, not a literature review research.

- Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods used compared to the research in this article. The difference lies in the fact that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].
- Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [33]. However, the previous research differs in the purpose and method compared to this article's research.
- Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of seriously Covid-19 patients with plasma proteomics [34]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

By referring to the elaboration of the most recent

previous related work by some researchers, the research carried out in this article has novelties that previous researchers have not studied. The novelty of the study lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

III. METHODOLOGY

This study applies two kinds of data mining methods or machine learning algorithms: RF and SVM. The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The development of the application program in this study uses the Python programming language.

The Data Mining Process for Covid-19 patient big data in this study uses CRISP-DM (Cross-Industry Standard Process for Data Mining) process; CRISP-DM is standard data mining. The CRISP-DM process comprises a six-stage [35]; see Figure 1 [36]. Figure 2 shows the data mining process in this study. This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

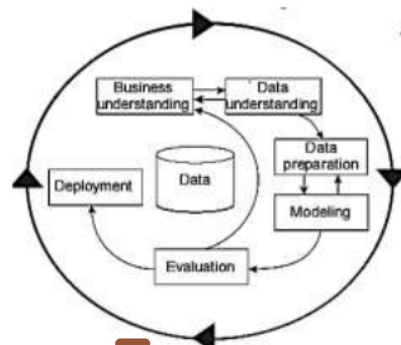


Figure 1. CRISP-DM Process

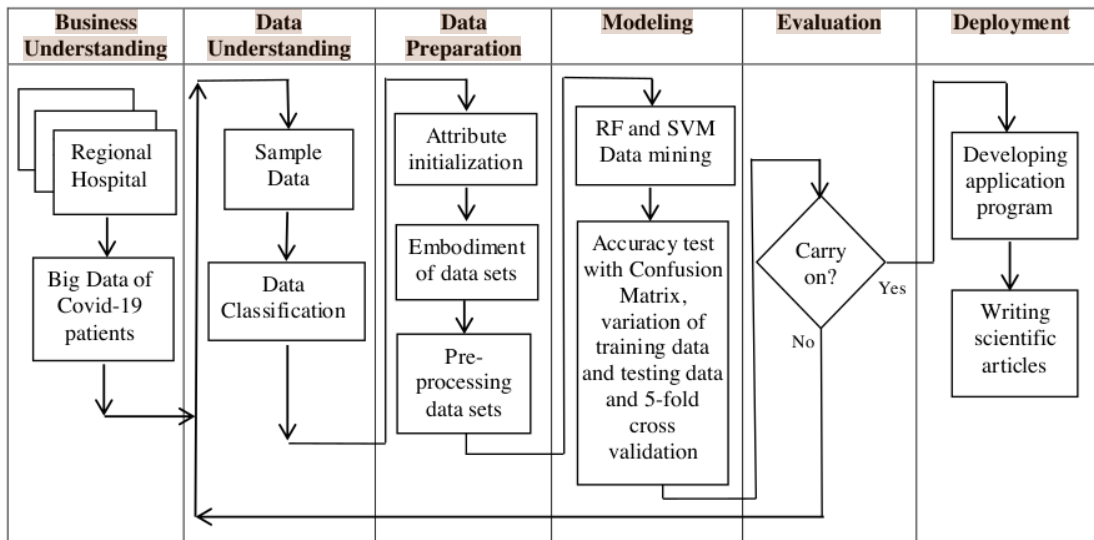


Figure 2. Data Mining Process of Covid-19 Patient Big Data

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected in the form of medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. The big data of Covid-19 patients obtained was 117. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 1 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE I. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses to others. In this study, the number of signs and symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 2).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or Gi where i = 1, 2, 3 ...12) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 3. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or

symptoms.

TABLE II. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE III. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	yes	no	no	No	No	No	Inpatient
2	yes	yes	no	Yes	No	No	Inpatient
3	yes	no	yes	No	No	No	Inpatient
4	yes	no	no	No	No	No	Inpatient
5	yes	no	no	No	No	No	Inpatient
6	yes	no	no	No	Yes	Yes	Inpatient
7	no	no	yes	No	No	No	Inpatient
...
...
114	no	no	no	No	No	Yes	Self-isolation
115	yes	no	no	No	Yes	No	Inpatient
116	yes	no	no	No	Yes	Yes	Inpatient
117	no	no	no	No	No	No	Inpatient

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1.

In addition, preprocessing the data set is also carried out on the class attribute, namely changing the class attribute with y and changing the value of the self-isolation class attribute with the number 1 and the *inpatient class attribute with the number -1, as shown in Table 4.

TABLE IV. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
...
...
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

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The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [37], which has their respective advantages in building application programs [38][39]. The application program built in this research uses the Python programming language to facilitate patient care status prediction.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

The formula of the SVM data mining method is: to form a linear equation from the training data; find the value of w and b, and the value of the classification decision is as follows.

$$S = ((x1, y1), \dots, (xl, yl)) \quad (1)$$

$$yi ((w \cdot xi) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_j^2 \quad (4)$$

$$Entropy(S) = \sum_i - p_i \times \log_2(p_i) \quad (5)$$

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

The format of the confusion matrix table is as shown in Table 5.

TABLE V. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-Validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation is helpful in assessing the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each group to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken obviously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the

training set (Dtrain). In the second fold, the second subset is the validate set the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 6. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE VI. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TEST DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% testing data and 90% training data and 10% testing data are as shown in Table 7.

TABLE VII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data		Accuracy		Precision		Recall	
(in %)		(in %)		(in %)		(in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

4.6 Deployment

One of the deployments in this research is in the form of making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 92%; In comparison, the SVM machine learning method has a prediction accuracy of 83% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Suggestions for further research are to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion is to conduct further research using other predictive data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first author, second, and so on as the fifth author.

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Abstract—Covid-19 disease spread throughout the world and caused death globally. Accordingly, researchers use many research methods to control the Covid-19 pandemic, one of which is getting more attention is the use of prediction methods. However, the problem in medical action is not easy to determine the treatment status. So it is not surprising that previous researchers emphasized that the most frequent mistakes were due to inaccuracies in medical decisions. Therefore, this study propose data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analytics or better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these hospitalized Covid-19 patients are receiving intensive medical care from doctors.

Errors in decision-making often occur because

decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [19][20][21]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [22]. For this reason, or that is why this study proposes a machine learning system model for decision-making solutions (predictions) of the treatment status of Covid-19 patients, both inpatient and self-isolation using data mining methods.

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [23]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting the new data with high accuracy [22].

Many Data Mining methods include K-Means,

Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out in order to be able to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [24] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [25] [26], as well as SVM algorithm has a widely known technique used for classification [26].

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some of the related works from the latest scientific articles compared with the work in this research article.

- Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [24]. This previous research is different from the research in the article on the research objectives and the object under study.
- Celestine Wendi et al. (2020) proposed the Random Forest model to predict disease severity of Covid-19 patients [27]. The difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and

SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.

- Chelvia Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [28]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, research in this article predicts the treatment status of Covid-19 patients.
- Based on patient clinical data, Boran Hao et al. (2020) developed a model to predict pneumonia severity and the level of care required by Covid-19 patients using statistical methods [29]. This previous research differs in the research purpose and way compared to the research in this article.
- Anthony Anggrawan et al. (2021) implemented machine learning to diagnose drug users and types of drug-using Forward Chaining and Certainty Factor methods [22]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.
- Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson distribution and the gamma distribution [30]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article.
- Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research objectives and methods used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. This article's research does not use the DT method but the RF and SVM methods.
- Sum Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [31]. This previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not using the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods.
- Ankit Mehrotra and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [32]. This previous research is a literature review study that concludes that the Data

Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from the research in this article because this article is an experimental study, not a literature review research.

- Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods used compared to the research in this article. The difference lies in the fact that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].
- Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [33]. However, the previous research differs in the purpose and method compared to this article's research.
- Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of seriously Covid-19 patients with plasma proteomics [34]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

By referring to the elaboration of the most recent

previous related work by some researchers, the research carried out in this article has novelties that previous researchers have not studied. The novelty of the study lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

III. METHODOLOGY

This study applies two kinds of data mining methods or machine learning algorithms: RF and SVM. The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The development of the application program in this study uses the Python programming language.

The Data Mining Process for Covid-19 patient big data in this study uses CRISP-DM (Cross-Industry Standard Process for Data Mining) process; CRISP-DM is standard data mining. The CRISP-DM process comprises a six-stage [35]; see Figure 1 [36]. Figure 2 shows the data mining process in this study. This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

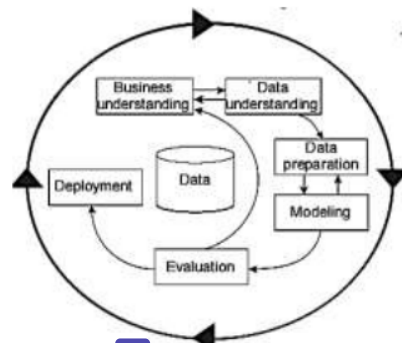


Figure 1. CRISP-DM Process

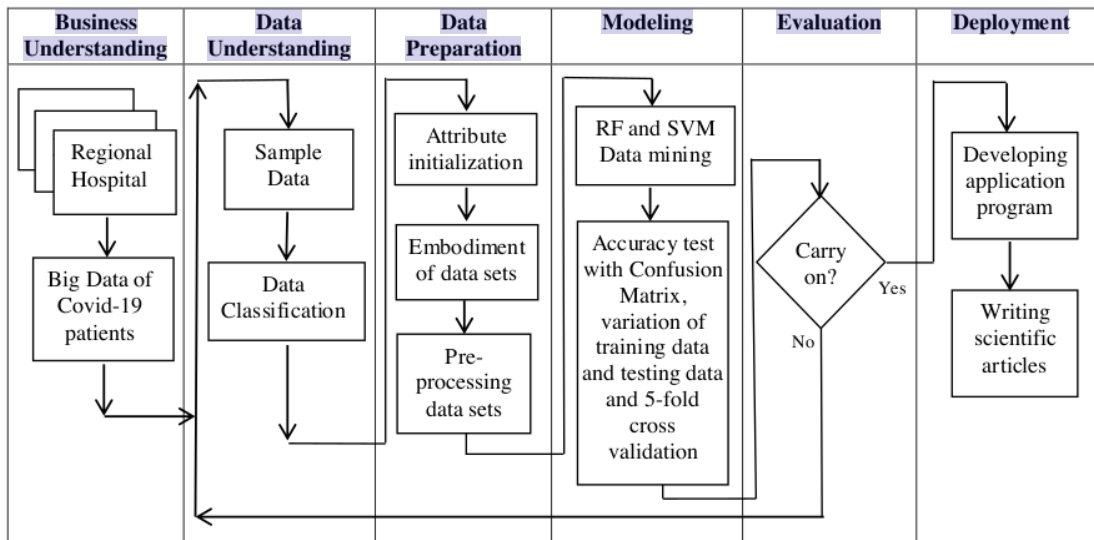


Figure 2. Data Mining Process of Covid-19 Patient Big Data

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected in the form of medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. The big data of Covid-19 patients obtained was 117. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 1 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE I. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses to others. In this study, the number of signs and symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 2).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or Gi where i = 1, 2, 3 ...12) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 3. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or

symptoms.

TABLE II. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE III. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	yes	no	no	No	No	No	Inpatient
2	yes	yes	no	Yes	No	No	Inpatient
3	yes	no	yes	No	No	No	Inpatient
4	yes	no	no	No	No	No	Inpatient
5	yes	no	no	No	No	No	Inpatient
6	yes	no	no	No	Yes	Yes	Inpatient
7	no	no	yes	No	No	No	Inpatient
...
...
114	no	no	no	No	No	Yes	Self-isolation
115	yes	no	no	No	Yes	No	Inpatient
116	yes	no	no	No	Yes	Yes	Inpatient
117	no	no	no	No	No	No	Inpatient

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1.

In addition, preprocessing the data set is also carried out on the class attribute, namely changing the class attribute with y and changing the value of the self-isolation class attribute with the number 1 and the *inpatient class attribute with the number -1, as shown in Table 4.

TABLE IV. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
...
...
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

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The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [37], which has their respective advantages in building application programs [38][39]. The application program built in this research uses the Python programming language to facilitate patient care status prediction.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

The formula of the SVM data mining method is: to form a linear equation from the training data; find the value of w and b, and the value of the classification decision is as follows.

$$S = ((x1, y1), \dots, (xl, yl)) \quad (1)$$

$$yi ((w \cdot xi) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum p_i^2 \quad (4)$$

$$Entropy(S) = \sum p_i \times \log_2(p_i) \quad (5)$$

4.4.3 Confusion matrix

This research uses a confusion matrix to measure performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

The format of the confusion matrix table is as shown in Table 5.

TABLE V. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-Validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation is helpful in assessing the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each group to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken obviously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the

training set (Dtrain). In the second fold, the second subset is the validate set the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 6. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE VI. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TEST DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

Further comparison of the accuracy and precision of the prediction system model with 40% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% testing data and 90% training data and 10% testing data are as shown in Table 7.

TABLE VII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)		Accuracy (in %)		Precision (in %)		Recall (in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

4.6 Deployment

One of the deployments in this research is in the form of making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 92%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Suggestions for further research are to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion is to conduct further research using other predictive data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first author, second, and so on as the fifth author.

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






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Abstract—Covid-19 disease spread throughout the world and caused death globally. Accordingly, researchers use many research methods to control the Covid-19 pandemic, one of which is getting more attention is the use of prediction methods. However, the problem in medical action is not easy to determine the treatment status. So it is not surprising that previous researchers emphasized that the most frequent mistakes were due to inaccuracies in medical decisions. Therefore, this study proposes a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analysis or better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these hospitalized Covid-19 patients are receiving intensive medical care from doctors.

Errors in decision-making often occur because

decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [19][20][21]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [22]. For this reason, or that is why this study proposes a machine learning system model for decision-making solutions (predictions) of the treatment status of Covid-19 patients, both inpatient and self-isolation using data mining methods.

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [23]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting the new data with high accuracy [22].

Many Data Mining methods include K-Means,

Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out in order to be able to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [24] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [25] [26], as well as SVM algorithm has a widely known technique used for classification [26].

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some of the related works from the latest scientific articles compared with the work in this research article.

- Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [24]. This previous research is different from the research in the article on the research objectives and the object under study.
- Celestine Iwendi et al. (2020) proposed the Random Forest model to predict disease severity of Covid-19 patients [27]. The difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.
- Chelvia Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [28]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, research in this article predicts the treatment status of Covid-19 patients.
- Based on patient clinical data, Boran Hao et al. (2020) developed a model to predict pneumonia severity and the level of care required by Covid-19 patients using statistical methods.[29]. This previous research differs in the research purpose and way compared to the research in this article.
- Anthony Anggrawan et al (2021) implemented machine learning to diagnose drug users and types of drug-using Ford Chaining and Certainty Factor methods [22]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.
- Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson distribution and the gamma distribution [30]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article.
- Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research objectives and methods used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. This article's research does not use the DT method but the RF and SVM methods.
- Sum Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [31]. This previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not using the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods.
- Ankit Mehrotra and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [32]. This previous research is a literature review study that concludes that the Data

Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from the research in this article because this article is an experimental study, not a literature review research.

- Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods used compared to the research in this article. The difference lies in the fact that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].
- Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [33]. However, the previous research differs in the purpose and method compared to this article's research.
- Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of seriously Covid-19 patients with plasma proteomics [34]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

By referring to the elaboration of the most recent

previous related work by some researchers, the research carried out in this article has novelties that previous researchers have not studied. The novelty of the study lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

III. METHODOLOGY

This study applies two kinds of data mining methods or machine learning algorithms: RF and SVM. The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The development of the application program in this study uses the Python programming language.

The Data Mining Process for Covid-19 patient big data in this study uses CRISP-DM (Cross-Industry Standard Process for Data Mining) process; CRISP-DM is standard data mining. The CRISP-DM process comprises a six-stage [35]; see Figure 1 [36]. Figure 2 shows the data mining process in this study. This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

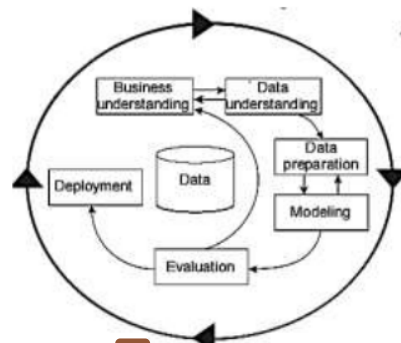


Figure 1. CRISP-DM Process

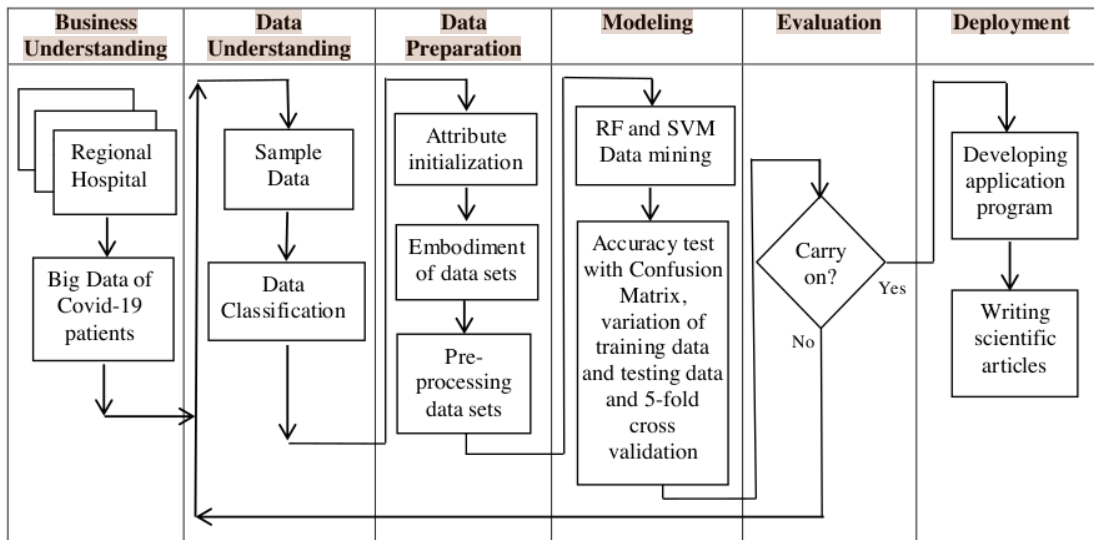


Figure 2. Data Mining Process of Covid-19 Patient Big Data

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected in the form of medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. The big data of Covid-19 patients obtained was 117. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 1 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE I. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses to others. In this study, the number of signs and symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 2).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or Gi where i = 1, 2, 3 ...12) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 3. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or

symptoms.

TABLE II. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE III. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	yes	no	no	No	No	No	Inpatient
2	yes	yes	no	Yes	No	No	Inpatient
3	yes	no	yes	No	No	No	Inpatient
4	yes	no	no	No	No	No	Inpatient
5	yes	no	no	No	No	No	Inpatient
6	yes	no	no	No	Yes	Yes	Inpatient
7	no	no	yes	No	No	No	Inpatient
...
...
114	no	no	no	No	No	Yes	Self-isolation
115	yes	no	no	No	Yes	No	Inpatient
116	yes	no	no	No	Yes	Yes	Inpatient
117	no	no	no	No	No	No	Inpatient

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1.

In addition, preprocessing the data set is also carried out on the class attribute, namely changing the class attribute with y and changing the value of the self-isolation class attribute with the number 1 and the *inpatient class attribute with the number -1, as shown in Table 4.

TABLE IV. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
...
...
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

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The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [37], which has their respective advantages in building application programs [38][39]. The application program built in this research uses the Python programming language to facilitate patient care status prediction.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

The formula of the SVM data mining method is: to form a linear equation from the training data; find the value of w and b, and the value of the classification decision is as follows.

$$S = ((x1, y1), \dots, (xl, yl)) \quad (1)$$

$$yi ((w \cdot xi) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_j^2 \quad (4)$$

$$Entropy(S) = \sum_i - p_i \times \log_2(p_i) \quad (5)$$

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

The format of the confusion matrix table is as shown in Table 5.

TABLE V. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-Validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation is helpful in assessing the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each group to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken obviously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the

training set (Dtrain). In the second fold, the second subset is the validate set the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 6. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE VI. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TEST DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% testing data and 90% training data and 10% testing data are as shown in Table 7.

TABLE VII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data		Accuracy		Precision		Recall	
(in %)		(in %)		(in %)		(in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

4.6 Deployment

One of the deployments in this research is in the form of making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 92%; In comparison, the SVM machine learning method has a prediction accuracy of 83% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Suggestions for further research are to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion is to conduct further research using other predictive data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first author, second, and so on as the fifth author.

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Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

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Abstract—Covid-19 has become a global pandemic that causes many deaths, so medical treatment for Covid-19 patients gets special attention, whether hospitalized or self-isolated. However, the problem in medical action is not easy, and the most frequent mistakes are due to inaccuracies in medical decision-making. Meanwhile, machine learning can predict with high accuracy. For that, or that's why this study aims to propose a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients. The implication is that RF machine learning can help/replace the role of medical experts in predicting the patient's care status.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not only because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analytics for better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

In essence, the care status for Covid-19 patients is self-isolation for patients with non-severe illness status and hospitalization for patients who are seriously ill and at critical risk or cause death. Hospitals or medical doctors take various ways to reduce the number of deaths of Covid-19 patients, including by regulating the status of services in hospitals. Expert doctors recommend that patients self-isolate or should be hospitalized after analyzing the patient's medical data. Determining the level of care for Covid-19 patients is a form of medical treatment or treatment for Covid-19 patients to get proper treatment or care.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18][19], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [20][21][22]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [23]. For this reason (or

why), this study's objective is to propose a machine learning system model for decision-making solutions (predictions) for the treatment status of Covid-19 patients, both inpatient and self-isolation, using data mining methods.

The implication is that the proposed machine learning system model can help and even replace the role of medical experts (specialist doctors) in making medical decisions for Covid-19 patients, whether hospitalization or self-isolation. That is, machine learning performs tasks like a medical specialist in deciding the results of the diagnosis of the nursing status of Covid-19 patients based on medical data of Covid-19 patients. Furthermore, machine learning can work tirelessly, time and place, and has intelligence like an expert, so it is not surprising that previous research confirms that intelligent machines can make superior decisions to experts because humans have a human error factor [24].

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [25]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting new data with high accuracy [23]. After all, predicting individuals with symptoms of being infected with Covid-19 mandates machine learning (application-based) and contributes to effectively isolating Covid-19 patients [26].

Big data demands large storage media [27]. However, big data is no longer traditionally processed [28]. Instead, today's big data processing relies on machines that can provide systematic results [29]. Big data storage is generally on a computer server with a large storage capacity. Still, some also make it happen by renting online cloud data storage services such as Amazon Simple Storage Service (Amazon S3) and Google Cloud [30]. Cloud facilitates cost-effective big data storage and analysis [30].

Big data processing techniques in data mining include several stages: target data, preprocessed data, data mining, and evaluation/analysis of mining results [30]. Target data and preprocessed data are the processes of extracting raw data from big data [30]. Target data is to select the required data (sample data) and classify data. The preprocessed data is to prepare data sets for data mining, including cleaning up incomplete data, duplicate data, and converting string data into numeric coded data. Finally, data mining and evaluation results extract hidden information by applying data mining methods suitable for the objectives and analyzing them.

Many Data Mining methods include K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative

Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [31] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [32] [33], as well as SVM algorithm has a widely known technique used for classification [33]. It is why this study uses SVM and RF to classify treatment status. Given that the SVM and RF machine learning algorithms are both popularly used by many researchers, the SVM and RF machine learning algorithms are the most appropriate combination used together in research, including research to classify and/or predict the treatment status of Covid-19 patients.

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some related works from the latest scientific articles compared with the work in this research article.

Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [31]. This previous research is different from the research in the article on the research objectives and the object under study. In the meantime, Celestine Iwendi et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [34]. The

difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.

Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [35]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, the research in this article predicts the treatment status of Covid-19 patients.

Based on patient clinical data, using statistical methods, Boran Hao et al. (2020) developed a model to predict pneumonia severity in Covid-19 patients using the Natural Language Processing tool [36]. However, this previous research differs in the research purpose and way compared to the research in this article. Meanwhile, Anthony Anggrawan et al. (2021) implemented machine learning to diagnose drug users and types of drug-using Forward Chaining and Certainty Factor methods [23]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.

Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson and Gamma distribution [37]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article. In the meantime, Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research objectives and techniques used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. This article's research does not use the DT method but uses the RF and SVM methods.

Soham Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [38]. However, this previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not utilizing the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods.

Whereas, Ankit Mehrotraa and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [39]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from this article's research because it is an experimental study, not a literature review research.

Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods compared to this article's research. The difference is that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].

Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [40]. However, the previous research differs in the purpose and way compared to this article's research. In contrast, Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of seriously ill Covid-19 patients with plasma proteomics [41]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

Table 1 shows a comparison between some of the most recent previous related work with the work carried out in this study. By referring to the elaboration of the most recent last related work by some researchers, the research carried out in this article has novelties (from the last research gap) that previous researchers have not studied. In essence, the gap in earlier research is that no one has researched machine learning models to predict the inpatient status or self-isolation of Covid-19 patients by involving RF and SVM algorithms. In addition, the 12 criteria used to indicate the treatment status of Covid-19 patients are entirely different from previous similar studies (as shown in Table 1 in the Criteria/Attributes column). So, the study's originality lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

TABLE I. COMPARISON OF THIS ARTICLE'S WORK WITH SOME PREVIOUS RELATED WORKS

Research by	Research methods			Criteria/Attributes		Research Object	Accuracy Test
	RF	SVM	ML	Number	Name		
Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) [31]	Yes	Yes	Yes	10	Coastal Aerosol, Blue, Green, Red, Vegetation Red Edge, NIR, Narrow NIR, Water vapor, SWIR-Cirrus, SWIR	Satellite remote sensing images	Yes
Celestine Iwendi et al. (2020) [34]	Yes	No	Yes	6	Symptom1, symptom2, symptom3, symptom4, symptom5, symptom6	Illness severity of Covid-19 patients	No
Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) [35]	Yes	Yes	Yes	9	Age, Body Mass Index (BMI), Glucose, Insulin, Homa, Leptin, Adiponectin, Resistin, MCP 1	Breast cancer	Yes
Boran Hao et al. (2020) [36]	No	No	No	10	Radiology Opacities, Respiratory Rate, Age, Fever Male, Albumin, Anion Gap, SpO2, LDH, Calcium	The severity of pneumonia in Covid-19 patients	Yes
Anthony Anggrawan et al. (2021) [23]	No	No	Yes	27	Out of breath, Anxious, Nausea, Diarrhea, Convulsions, Easily angry, Depression, Sleep patterns change, Sweating, Chills, Shaking, Insomnia, Fast heart rate, Blood pressure rises, Difficult to focus, Difficult to rest, Weight loss, Dry mouth, Blurred vision, Changed skin color, Constipation, Stomachache, Drowsiness, Itching, Difficulty urinating, Mood swings, Dizziness	Drug users and types of drug-using	Yes
Hongwei Zhao et al. (2021) [37]	No	No	No	0	-	Number of cases of Covid-19 patients	No
Bassam Mahboub et al. (2021) [8]	No	No	Yes	5	Age, Gender, Nationality, Blood group, BMI	The treatment period for Covid-19 patient	Yes
Soham Guhathakurata et al. (2021) [38]	No	Yes	Yes	8	Temp, Breathing rate, Hypertension, Heartbeat rate (HBR), Acute respiratory disease syndrome (ARDS), Chest pain, Heart disease, Cough with sputum (CWS)	Predicting whether patients are infected with Covid-19 or not	No
Ankit Mehrotraa and Reeti Agarwal (2021) [39]	No	No	No	0	-	Discussing the data mining method's role in the Covid-19 pandemic	No
Pratiyush Guleria et al. (2022) [14]	No	Yes	Yes	0	-	The death rate of Covid-19 patient	Yes
Anthony Anggrawan, et al. (2022) [40]	No	No	Yes	6	Achievement index, achievement points, recommendation, organizational activity, semester level, and completeness of documents	Scholarship recipient	Yes
Vadim Demichev et al. (2022) [41]	No	No	No	0	-	Optimization of treatment for Covid-19 patients	Yes
Our research	Yes	Yes	Yes	12	Pneumonia, ARDS, CHF, AKI, CAD, Dyspnea, NSTEMI, ADHF, HHD, Febris, Anosmia, Ageusia	Care status of Covid-19 patients, whether inpatient or self-isolation	Yes

Note: ML = Machine Learning

III. METHODOLOGY

This study applies two data mining methods or machine learning algorithms: RF and SVM. The big data is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The big data source used in this study is primary data from patient medical records/documents. The attributes of the patient's disease symptoms and care status classes amount to thousands of patient medical record data. Patient datasets containing non-Covid-19 and duplicate and incomplete Covid-19 patient data are removed, so only data is left as a dataset for data mining processes. Medical record data of disease symptoms obtained

from string data is then converted into numeric data. The development of the application program in this study uses the Python computer programming language.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods. The data mining process in this research uses CRISP-DM (Cross-Industry Standard Process for Data Mining). CRISP-DM is a standard data mining process. The process in CRISP-DM comprises a six-stage [42], as shown in Figure 1 [43]. Figure 2 shows the process carried out at each stage of CRISP-DM.

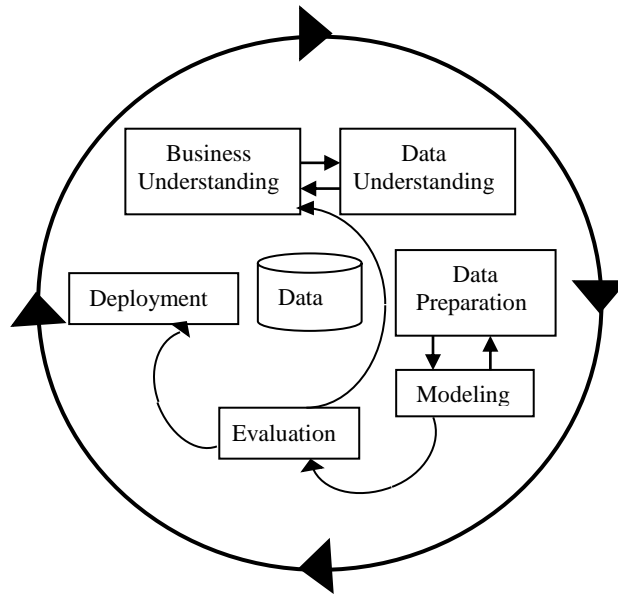


Figure 1. The CRISP-DM Process

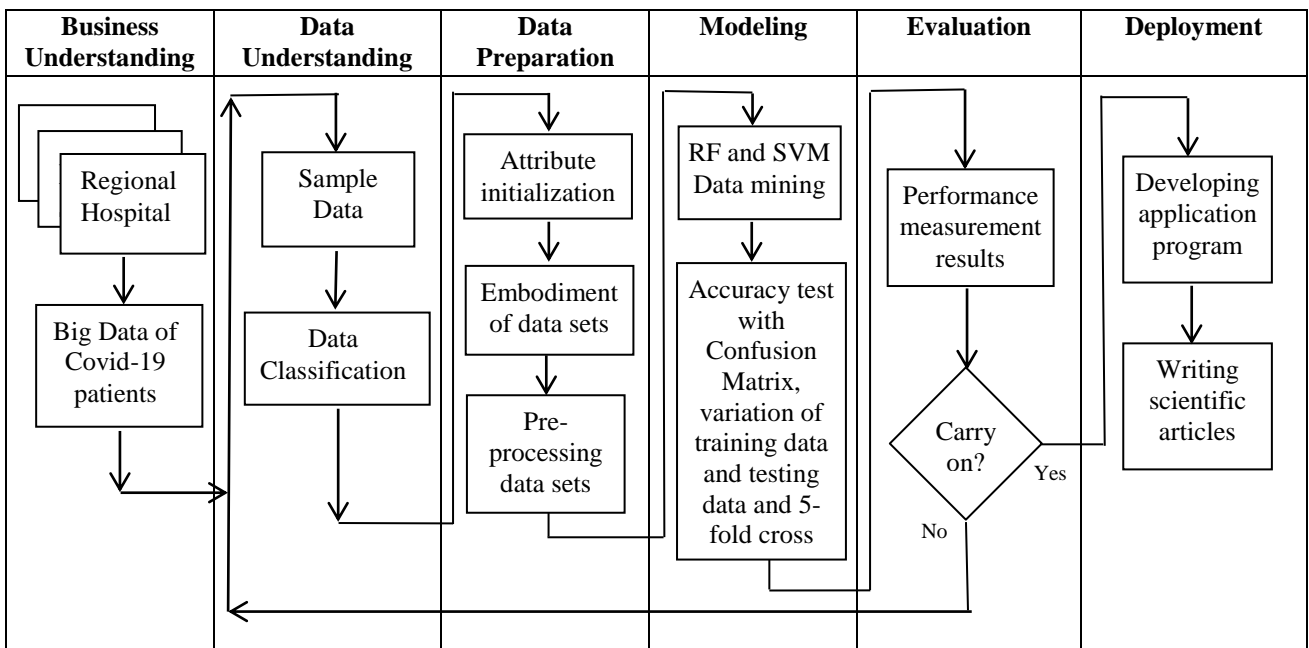


Figure 2. Data Mining Process of Covid-19 Patient Big Data

In Figure 2, business understanding is the stage of sorting out thousands of hospital patient medical data to collect the required patient data. The next stage is understanding the data collected as representative data for Covid-19 patients. This Covid-19 patient data classifies the patient's signs and symptoms and treatment status, which needs further processing at the next stage. The next stage is the data preparation stage, which essentially determines the attributes of the names of signs and symptoms of Covid-19 patients. The embodiment of the dataset containing knowledge according to treatment status refers to the signs and symptoms that the patient has (marked with Yes) or does not have (marked with No). The next thing to do is preprocess the dataset, changing the category value of the symptom attribute and the class attribute with the number 1 and the number -1. The process of extracting raw data obtained in the previous process stage is data that is further processed by data mining methods or used as learning machine learning data at the modeling stage. So that machine learning can predict. The next stage is the evaluation stage, namely, knowing the predictive reliability of the data mining or machine learning method. Then the last stage is the deployment stage to disseminate research results so that they are helpful for implementation by various parties, especially hospitals and other professionals, in the form of developing application programs and scientific articles.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected from medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. There are thousands of data on Covid-19 patients. The patient dataset containing incomplete data and non-covid-19 was omitted or ignored. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 2 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE II. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses. There were 117 patients with Covid-19 who had a different diagnosis from the others. In this study, the number of signs and symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 3).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or G_i where $i = 1, 2, 3 \dots 12$) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 4. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or symptoms.

Furthermore, the preprocessing of the data set is done by changing the G_i with x_i and the G_i Yes attribute value with the number 1 while the G_i No attribute with the number -1. In addition, dataset preprocessing is also carried out on class attributes, namely changing the independent isolation class attribute category with the number 1 and the inpatient class attribute category with the number -1, as shown in Table 5.

TABLE III. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE IV. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	Yes	No	No	No	No	No	Inpatient
2	Yes	Yes	No	Yes	No	No	Inpatient
3	Yes	No	Yes	No	No	No	Inpatient
4	Yes	No	No	No	No	No	Inpatient
5	Yes	No	No	No	No	No	Inpatient
6	Yes	No	No	No	Yes	Yes	Inpatient
7	No	No	Yes	No	No	No	Inpatient
..
..
114	No	No	No	No	No	Yes	Self-isolation
115	Yes	No	No	No	Yes	No	Inpatient
116	Yes	No	No	No	Yes	Yes	Inpatient
117	No	No	No	No	No	No	Inpatient

TABLE V. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
..
..
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [44], which has their respective advantages in building application programs [45][46]. The application program built in this research uses the Python programming language to facilitate patient care status prediction. The hyperparameter used in the SVM method is kernel = linear and Probability = True. Meanwhile, the hyperparameter used for the RF hyperparameter method is n_estimators = 100. The n_estimators is the number of trees in RF.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

In SVM, there are two implementation models: mathematical programming techniques and kernel functions. This study applies kernel functions and focuses on classifying two categories of class attributes. The class attribute is a treatment for $y_i = +1, -1$. The formula of the SVM data mining method is: (a) to form a linear equation from the training data; (b) find the value of w and b, and (c) the value of the classification decision is as follows.

$$S = ((x1, y1), \dots, (xl, yl)) \quad (1)$$

$$y_i ((w \cdot x_i) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

RF uses bootstrap samples from training data to create a tree from a randomly selected subset. The chosen predictor is a candidate for splitting the decision tree. The results of the category predictions from the treatment class based on the results of the highest voting were chosen as the final prediction results. The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_{i=1}^c p_i^2 \quad (4)$$

$$Entropy(S) = \sum_{i=1}^c -p_i \times \log_2(p_i) \quad (5)$$

Description:

S = Set of cases

p_i = the proportion of case i to the Set of cases

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and Recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

Accuracy states the closeness of the measurement results to the actual value, while Precision shows how close the difference in the measurement results is on repeated measurements. On the other hand, Recall states the level of success in retrieving information. Precision and Recall are necessary because Precision denotes a measure of quality, and Recall denotes a measure of quantity.

Measurement of accuracy is based on the ratio between the correct predictions (positive and negative) with the overall data. In contrast, precision measurements are based on the percentage of true positive predictions compared to overall positive predicted outcomes. Meanwhile, the recall measurement is based on the ratio of true positive predictions compared to the general actual positive data.

The format of the confusion matrix table is as shown in Table 6.

TABLE VI. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-Validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation helps assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold value. In the

performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each subgroup to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the training set (Dtrain). In the second fold, the second subset is the validate set, the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 7 and Figure 3. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE VII. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TESTING DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

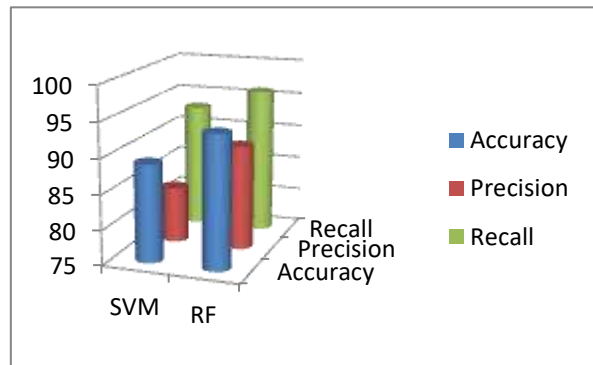


Figure 3. System Model Performance Testing with 85% of Training Data and 15% of Testing Data

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% data 90% testing and training data and 10% testing data are as shown in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)		Accuracy (in %)		Precision (in %)		Recall (in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix. 333

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 9 and Figure 4.

TABLE IX. PREDICTION PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

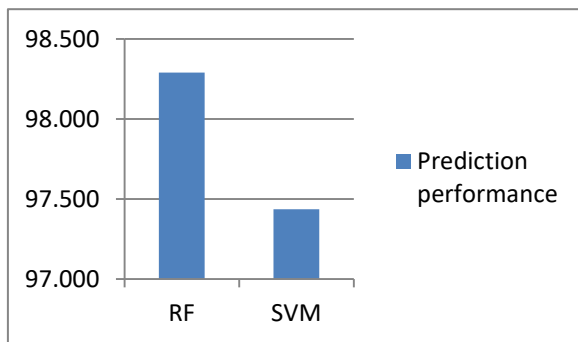


Figure 4. Predictive performance testing using K-fold cross-validation

4.6 Deployment

One of the deployments in this research is making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Thus, the results obtained can be developed and become the knowledge of many parties as a responsibility for the correctness of the effects of research carried out as professional researchers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 90%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%. The research result implication is that RF machine learning can help or replace the role of medical personnel in predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, with high accuracy.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Further research needs to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion for future study is: to conduct further research using other data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients and various other diseases, to build a system that not only predicts but also performs clustering, association, and estimates of various other fields of science, including patients' care status, with a combination of machine learning and the Internet of Things.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first, second, and so on as the fifth author.

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Correction Note

Manuscript ID: JAIT-5790

Title: Machine Learning to Predict the Treatment Status of Covid-19 Patients

Dear Reviewers

We have made improvements to our manuscript following the instructions and suggestions of the reviewers. About which parts we have revised and added writing, it is as explained below:

	Reviewer's Comments (Reviewer A)	The Change/Corrections made (the change is marked with a yellow highlight)	In Page & paragraph
1	Abstract: is vague and does not serve its purpose. Abstract should introduce the domain, discuss the problem statement, present the objective, discuss the possible outcome, and finally present the contribution to the research domain	<p>Abstract Contents after revision</p> <p>Covid-19 has become a global pandemic that causes many deaths, so medical treatment for Covid-19 patients gets special attention, whether hospitalized or self-isolated. However, the problem in medical action is not easy, and the most frequent mistakes are due to inaccuracies in medical decision-making. Meanwhile, machine learning can predict with high accuracy. For that, or that's why this study aims to propose a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients. The implication is that RF machine learning can help/replace the role of medical experts in predicting the patient's care status.</p>	On page 1
2	Introduction: this section is again vague. The background on Covid-19 is good but it is expected to discuss more from the research perspective. What are the research implications of the Covid-19 pandemic? How is artificial intelligence contributing to controlling the pandemic? What is the research gap you have identified? Why are you	<p>Following the reviewer's suggestion, the authors have added writing related to the reviewer's direction in the Introduction subsection and Related works subsection as follows:</p> <p>* In essence, the care status for Covid-19 patients is self-isolation for patients with non-severe illness status and hospitalization for patients who are seriously ill and at critical risk or cause death. Hospitals or medical doctors take various ways to reduce the number of deaths of Covid-19 patients, including by regulating the status of services in hospitals. Expert doctors recommend that patients self-isolate or should be hospitalized after</p>	On page 2, 2 nd paragraph, 2 nd column, in the Introduction subsection

	<p>proposing this work? All of these above questions should be answered within the introduction and related works section.</p>	<p>analyzing the patient's medical data. Determining the level of care for Covid-19 patients is a form of medical treatment or treatment for Covid-19 patients to get proper treatment or care.</p> <p>* For this reason, this study's objective is to propose a machine learning system model for decision-making solutions (predictions) for the treatment status of Covid-19 patients, both inpatient and self-isolation, using data mining methods.</p> <p>The implication is that the proposed machine learning system model can help and even replace the role of medical experts (specialist doctors) in making medical decisions for Covid-19 patients, whether hospitalization or self-isolation. That is, machine learning performs tasks like a medical specialist in deciding the results of the diagnosis of the nursing status of Covid-19 patients based on medical data of Covid-19 patients. However, machine learning can work tirelessly, time and place, and has intelligence like an expert, so it is not surprising that previous research confirms that intelligent machines can make superior decisions to experts because humans have a human error factor [23].</p> <p>* After all, predicting individuals with symptoms of being infected with Covid-19 mandates machine learning (application-based) and contributes to effectively isolating Covid-19 patients [25].</p> <p>* By referring to the elaboration of the most recent previous related work by some researchers, the research carried out in this article has novelties (from the last research gap) that previous researchers have not studied. In essence, the gap in earlier research is that no one has researched machine learning models to predict the inpatient status or self-isolation of Covid-19 patients by involving RF and SVM algorithms.</p>	<p>Page 2, 1st paragraph, 1st column, in the Introduction subsection</p> <p>Page 2, 2nd paragraph, 1st column, in the Introduction subsection</p> <p>Page 2, 3rd paragraph, 1st column, in the Introduction subsection</p> <p>Page 3, 8th paragraph, 2nd column, in the Related Works subsection</p>
3	<p>Related works: This section summarizes a few research papers on RF and SVM however, there is no real analysis done. Related work study is to identify the relevance of the proposed work and to identify the research gap. I suggest the authors to add a table comparing the state-of-the-art</p>	<p>The authors have added Table 1. Comparison of This Article's Work With Some Previous Related Works in the Related Works subsection following the reviewers' suggestions</p> <p>The authors add sentences related to table 1 as follows: Table 1 shows a comparison between some of the most recent previous related work with the work carried out in this study</p>	<p>Page 4, in the Related Works subsection</p> <p>Page 3, 8th paragraph, 2nd column, in the Related Works subsection</p>

	<p>research on covid-19 prediction with machine learning.</p>		
<p>4</p>	<p>a. Methodology: is very weak. Generally, for a research paper, a methodology paper should be given more importance. It should clearly explain the dataset, introduce the methodology used, and discuss the novelty and algorithms if any.</p> <p>b. I felt the subsections belonging to results and discussions should be added to the methodology section and more detailed information should be added;</p> <p>c. Figure 1 is not readable and of poor image quality;</p> <p>d. SVM and RF hyperparameter details are not discussed. More details of SVM and RF should be given;</p> <p>e. Confusion matrix format is presented but the result is not plotted;</p> <p>f. The need for a recall, and precision is not discussed. The specificity result should be reported which is missing;</p> <p>g. Deployment details should be justified. How scientific article writing can be considered deployment?</p>	<p>The authors have added descriptions to the methodology subsections as follows: This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.</p> <p>a. The data mining process in this research uses CRISP-DM (Cross-Industry Standard Process for Data Mining). CRISP-DM is a standard data mining process. The process in CRISP-DM comprises a six-stage [41], as shown in Figure 1 [42]. Meanwhile, figure 2 shows the process carried out at each stage of CRISP-DM.</p> <p>b. Figure 2 shows the process carried out at each stage of CRISP-DM. In figure 2, Business understanding is the stage of sorting out thousands of hospital patient medical data to collect the required patient data. The next stage is understanding the data collected as representative data for Covid-19 patients. This Covid-19 patient data classifies the patient's signs and symptoms and treatment status, which needs further processing at the next stage. The next stage is the data preparation stage, which essentially determines the attributes of the names of signs and symptoms of Covid-19 patients. The embodiment of the dataset containing knowledge according to treatment status refers to the signs and symptoms that the patient has (marked with Yes) or does not have (marked with No). The next thing to do is preprocess the dataset, changing the category value of the symptom attribute and the class attribute with the number 1 and the number -1. The process of extracting raw data obtained in the previous process stage is data that is further processed by data mining methods or used as learning machine learning data at the modeling stage. So that machine learning can predict. The next stage is the evaluation stage, namely, knowing the predictive reliability of the data mining or machine learning method. Then the last stage is the deployment stage to disseminate research results so that they are helpful for implementation by various parties, especially hospitals and other professionals, in the form of developing application programs and scientific articles.</p> <p>c. The author has improved Figure 1 so that the image quality is legible and clear.</p> <p>d. The hyperparameter used in the SVM method is kernel = linear and probability = True; Meanwhile the hyperparameter used for the RF hyperparameter method is n_estimators = 100. n_estimators is the number of trees in RF.</p>	<p>Page 5, 2nd paragraph, 2nd column, in the methodology subsection</p> <p>Page 5, 2nd paragraph, 2nd column & page 6, 1st paragraph, 1st column</p> <p>On page 5</p> <p>On page 7, 2nd column, paragraph 4.</p>

		<p>e. Regarding the presentation of the confusion matrix format presented, the author has added the results in Figure 3 and Figure 4 (in Result and Discussion subsection).</p> <p>f. Accuracy states the closeness of the measurement results to the actual value, while Precision shows how close the difference in the measurement results is on repeated measurements. On the other hand, Recall states the level of success in retrieving information. Precision and Recall are necessary because Precision denotes a measure of quality, and Recall denotes a measure of quantity.</p> <p>g. One of the deployments in this research is making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Thus, the results obtained can be developed and become the knowledge of many parties as a responsibility for the correctness of the effects of research carried out as professional researchers.</p>	<p>Figure 3 on page 8 and Figure 4 on page 9</p> <p>On page 8, paragraph 3, 1st column</p> <p>On page 9, in subheading 4.6 Deployment</p>
5	<p>Future directions is weak. Should be more clear based on the findings and extension of the work.</p>	<p>As per the reviewer's suggestions, the authors have added the following description (text highlighted in yellow in the Conclusion subsection) as follows: Another suggestion for future study is: to conduct further research using other data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients and various other diseases, to build a system that not only predicts but also performs clustering, association, and estimates of various other fields of science, including patients' care status, with a combination of machine learning and the Internet of Things.</p>	<p>On page 9, 2nd column, paragraph 3, in Conclusion subsection</p>
6	<p>In many places, the authors have mentioned big data. When a bigdata is involved the processing and storing techniques of bigdata should also be discussed. Authors must clearly define the type of data they are using in the work.</p>	<p>In accordance with the reviewers' suggestions, the authors have added the following description to the manuscript: Big data demands large storage media [26]. However, big data is no longer traditionally processed [27]. Instead, today's big data processing relies on machines that can provide systematic results [28]. Big data storage is generally on a computer server with a large storage capacity. Still, some also make it happen by renting online cloud data storage services such as Amazon Simple Storage Service (Amazon S3) and Google Cloud [29]. Cloud facilitates cost-effective big data storage and analysis [29]. Big data processing techniques in data mining include several stages: target data, preprocessed data, data mining, and evaluation/analysis of mining results [29]. Target data and preprocessed data are the processes of extracting raw data from big data [29]. Target data is to select the required data (sample data) and classify data. The preprocessed data is to prepare data sets for data mining, including cleaning up incomplete data, duplicate data, and converting string data into numeric coded data. Finally, data mining and evaluation results extract hidden information by applying data mining methods suitable for the objectives and analyzing them.</p>	<p>On page 2, 1st column, paragraph 4 and 5, in Introduction subsection</p>

		Medical record data of disease symptoms obtained in the form of string data is then converted into numeric data.	On page 5, 1 st and 2 nd columns, in methodology subsection
7	<p>There were many typos, grammatical and colloquial language found. This should be avoided.</p> <p>Graphical representation is missing. Authors should consider adding more graphical representations of results.</p>	<p>The author has revised according to reviewers' suggestions on many parts of the manuscript (the change is marked with a light blue highlight).</p> <p>For example:</p> <ul style="list-style-type: none"> * The sentence "The data collected in the form of medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. " in the Result and Discussion subsection, it is replaced with the sentence "The data collected from medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified.". The phrase "in the form of" is replaced with "from". * the sentence "..... and a precision of 92%; ..." in the Conclusion subsection on lines 8 and 9, has been changed to "and a precision of 90%; ...". <p>As per reviewers' suggestions, the authors have added figure 2 and figure 4 charts.</p>	<p>On page 6, 1st column, in Result and Discussion subsection, rows 3 to 7.</p> <p>On page 9, 2nd column, in conclusion subsection, paragraph 1.</p>
8			
	Reviewer's Comments (Reviewer B)	Corrections made (the change is marked with a yellow highlight)	In Page & paragraph
1	In section II. Related works, the review of past work or studies by other researchers should be written in a paragraph. Currently, the authors used bullet style to describe each study. Then, a summary of the description should be described in table form. this would be easier for the reader to compare the previous studies with what has been proposed by the authors. Perhaps the table should provide details about the study, the authors, the limitation, etc.	The authors have followed the reviewer's suggestion of changing the related works review (on page 2 to 3 in the Related works subsection) in paragraphs instead of bullet style. Then, have summarized the description in table 1 (on page 4 in the Related works subsection).	On page 2 to 4, in Related Works subsection
2	Background of machine learning algorithms: RF and SVM should be provided and described in the manuscript.	The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [30] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for	On page 2, 2 nd column, paragraph 1, in Introduction

		classification [31] [32], as well as SVM algorithm has a widely known technique used for classification [32]. It is why this study uses SVM and RF to classify treatment status.	subsection
3	The review of other machine learning algorithms also should be provided, and the author should conclude or justify why RF and SVM are a good combination for this study.	Given that the SVM and RF machine learning algorithms are both popularly used by many researchers, the SVM and RF machine learning algorithms are the most appropriate combination used together in research, including research to classify and/or predict the treatment status of Covid-19 patients.	On page 2, 2 nd column, paragraph 1, in Introduction subsection
4	What kind of data was used in this study? Which source? Please cite and justify.	The author has justified the kind of data used in this study (in the methodology subsection) according to the reviewers' suggestions as follows: The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The big data source used in this study is primary data from patient medical records/documents.	On page 5, 1 st column, in methodology subsection
5	The authors claimed that this study involved big data. But in section IV Result and Discussion, the data used in this study is 117. The data is too small, thus contradicting the previous statement in the abstract and introduction.	The authors have added a description in the methodology subsection as follows: The attributes of the patient's disease symptoms and care status classes amount to thousands of patient medical record data. Patient datasets containing non-Covid-19 as well as duplicate and incomplete Covid-19 patient data are removed, so that only data is left as a dataset for data mining processes.	On page 5, 1 st paragraph, in the methodology subsection
6	In section 4.4.1, a detailed description of the SVM data mining method is not provided. The authors only provide general step but did not detail out the process.	The authors have added a description as follows: In SVM, there are two implementation models: mathematical programming techniques and kernel functions. This study applies kernel functions and focuses on classifying two categories of class attributes. The class attribute is a treatment for $y_i = +1, -1$.	On page 7, 2 nd column, in Result and Discussion subsection
7	Section 4.4.2, a detailed description of the RF data mining method is not provided. The authors only provide general step but did not detail out the process.	The authors have added a description as follows: RF uses bootstrap samples from training data to create a tree from a randomly selected subset. The chosen predictor is a candidate for splitting the decision tree. The results of the category predictions from the treatment class based on the results of the highest voting were chosen as the final prediction results. The formula for the RF data mining method is the Gini criterion and the Entropy criterion: $Gini = 1 - \sum_{i=1}^c p_i^2 \quad (4)$ $Entropy(S) = \sum_{i=1}^c -p_i \times \log_2(p_i) \quad (5)$ Description: S = Set of cases p_i = the proportion of case i to the Set of cases	On page 7, 2 nd column, in Result and Discussion subsection
8	The accuracy, Precision and Recall factor is based on what? It is unclear how you measure these factors.	The authors have added a description as follows: Measurement of accuracy is based on the ratio between the correct predictions (positive and negative) with the overall data. In contrast, precision measurements are based on the percentage of true positive predictions compared to overall	On page 8, column 1, paragraph 5, in Result and Discussion

		positive predicted outcomes. Meanwhile, the recall measurement is based on the ratio of true positive predictions compared to the general actual positive data.	subsection
9	In conclusion, this manuscript is basically comparing two methods of machine learning; RF and SVM. The title did not reflect this. The title should be revised. Example Comparative analysis of The current title could be misleading.	Authors: The manuscript title has been revised to Comparative analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients. Thanks you	

Hopefully, what we have done fulfills the wishes of the reviewers.
Thank you very much. God bless you.

Sincerely yours
Authors

Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

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Abstract—Covid-19 has become a global pandemic that causes many deaths, so medical treatment for Covid-19 patients gets special attention, whether hospitalized or self-isolated. However, the problem in medical action is not easy, and the most frequent mistakes are due to inaccuracies in medical decision-making. Meanwhile, machine learning can predict with high accuracy. For that, or that's why this study aims to propose a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients. The implication is that RF machine learning can help/replace the role of medical experts in predicting the patient's care status.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not only because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analytics for better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

In essence, the care status for Covid-19 patients is self-isolation for patients with non-severe illness status and hospitalization for patients who are seriously ill and at critical risk or cause death. Hospitals or medical doctors take various ways to reduce the number of deaths of Covid-19 patients, including by regulating the status of services in hospitals. Expert doctors recommend that patients self-isolate or should be hospitalized after analyzing the patient's medical data. Determining the level of care for Covid-19 patients is a form of medical treatment or treatment for Covid-19 patients to get proper treatment or care.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18][19], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [20][21][22]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [23]. For this reason (or

why), this study's objective is to propose a machine learning system model for decision-making solutions (predictions) for the treatment status of Covid-19 patients, both inpatient and self-isolation, using data mining methods.

The implication is that the proposed machine learning system model can help and even replace the role of medical experts (specialist doctors) in making medical decisions for Covid-19 patients, whether hospitalization or self-isolation. That is, machine learning performs tasks like a medical specialist in deciding the results of the diagnosis of the nursing status of Covid-19 patients based on medical data of Covid-19 patients. Furthermore, machine learning can work tirelessly, time and place, and has intelligence like an expert, so it is not surprising that previous research confirms that intelligent machines can make superior decisions to experts because humans have a human error factor [24].

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [25]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting new data with high accuracy [23]. After all, predicting individuals with symptoms of being infected with Covid-19 mandates machine learning (application-based) and contributes to effectively isolating Covid-19 patients [26].

Big data demands large storage media [27]. However, big data is no longer traditionally processed [28]. Instead, today's big data processing relies on machines that can provide systematic results [29]. Big data storage is generally on a computer server with a large storage capacity. Still, some also make it happen by renting online cloud data storage services such as Amazon Simple Storage Service (Amazon S3) and Google Cloud [30]. Cloud facilitates cost-effective big data storage and analysis [30].

Big data processing techniques in data mining include several stages: target data, preprocessed data, data mining, and evaluation/analysis of mining results [30]. Target data and preprocessed data are the processes of extracting raw data from big data [30]. Target data is to select the required data (sample data) and classify data. The preprocessed data is to prepare data sets for data mining, including cleaning up incomplete data, duplicate data, and converting string data into numeric coded data. Finally, data mining and evaluation results extract hidden information by applying data mining methods suitable for the objectives and analyzing them.

Many Data Mining methods include K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative

Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [31] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [32] [33], as well as SVM algorithm has a widely known technique used for classification [33]. It is why this study uses SVM and RF to classify treatment status. Given that the SVM and RF machine learning algorithms are both popularly used by many researchers, the SVM and RF machine learning algorithms are the most appropriate combination used together in research, including research to classify and/or predict the treatment status of Covid-19 patients.

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some related works from the latest scientific articles compared with the work in this research article.

Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [31]. This previous research is different from the research in the article on the research objectives and the object under study. In the meantime, Celestine Iwendi et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [34]. The

difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.

Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [35]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, the research in this article predicts the treatment status of Covid-19 patients.

Based on patient clinical data, using statistical methods, Boran Hao et al. (2020) developed a model to predict pneumonia severity in Covid-19 patients using the Natural Language Processing tool [36]. However, this previous research differs in the research purpose and way compared to the research in this article. Meanwhile, Anthony Anggrawan et al. (2021) implemented machine learning to diagnose drug users and types of drug-using Forward Chaining and Certainty Factor methods [23]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.

Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson and Gamma distribution [37]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article. In the meantime, Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research objectives and techniques used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. This article's research does not use the DT method but uses the RF and SVM methods.

Soham Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [38]. However, this previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not utilizing the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods.

Whereas, Ankit Mehrotraa and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [39]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from this article's research because it is an experimental study, not a literature review research.

Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods compared to this article's research. The difference is that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].

Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [40]. However, the previous research differs in the purpose and way compared to this article's research. In contrast, Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of seriously ill Covid-19 patients with plasma proteomics [41]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

Table 1 shows a comparison between some of the most recent previous related work with the work carried out in this study. By referring to the elaboration of the most recent last related work by some researchers, the research carried out in this article has novelties (from the last research gap) that previous researchers have not studied. In essence, the gap in earlier research is that no one has researched machine learning models to predict the inpatient status or self-isolation of Covid-19 patients by involving RF and SVM algorithms. In addition, the 12 criteria used to indicate the treatment status of Covid-19 patients are entirely different from previous similar studies (as shown in Table 1 in the Criteria/Attributes column). So, the study's originality lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

TABLE I. COMPARISON OF THIS ARTICLE'S WORK WITH SOME PREVIOUS RELATED WORKS

Research by	Research methods			Criteria/Attributes		Research Object	Accuracy Test
	RF	SVM	ML	Number	Name		
Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) [31]	Yes	Yes	Yes	10	Coastal Aerosol, Blue, Green, Red, Vegetation Red Edge, NIR, Narrow NIR, Water vapor, SWIR-Cirrus, SWIR	Satellite remote sensing images	Yes
Celestine Iwendi et al. (2020) [34]	Yes	No	Yes	6	Symptom1, symptom2, symptom3, symptom4, symptom5, symptom6	Illness severity of Covid-19 patients	No
Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) [35]	Yes	Yes	Yes	9	Age, Body Mass Index (BMI), Glucose, Insulin, Homa, Leptin, Adiponectin, Resistin, MCP 1	Breast cancer	Yes
Boran Hao et al. (2020) [36]	No	No	No	10	Radiology Opacities, Respiratory Rate, Age, Fever Male, Albumin, Anion Gap, SpO2, LDH, Calcium	The severity of pneumonia in Covid-19 patients	Yes
Anthony Anggrawan et al. (2021) [23]	No	No	Yes	27	Out of breath, Anxious, Nausea, Diarrhea, Convulsions, Easily angry, Depression, Sleep patterns change, Sweating, Chills, Shaking, Insomnia, Fast heart rate, Blood pressure rises, Difficult to focus, Difficult to rest, Weight loss, Dry mouth, Blurred vision, Changed skin color, Constipation, Stomachache, Drowsiness, Itching, Difficulty urinating, Mood swings, Dizziness	Drug users and types of drug-using	Yes
Hongwei Zhao et al. (2021) [37]	No	No	No	0	-	Number of cases of Covid-19 patients	No
Bassam Mahboub et al. (2021) [8]	No	No	Yes	5	Age, Gender, Nationality, Blood group, BMI	The treatment period for Covid-19 patient	Yes
Soham Guhathakurata et al. (2021) [38]	No	Yes	Yes	8	Temp, Breathing rate, Hypertension, Heartbeat rate (HBR), Acute respiratory disease syndrome (ARDS), Chest pain, Heart disease, Cough with sputum (CWS)	Predicting whether patients are infected with Covid-19 or not	No
Ankit Mehrotraa and Reeti Agarwal (2021) [39]	No	No	No	0	-	Discussing the data mining method's role in the Covid-19 pandemic	No
Pratiyush Guleria et al. (2022) [14]	No	Yes	Yes	0	-	The death rate of Covid-19 patient	Yes
Anthony Anggrawan, et al. (2022) [40]	No	No	Yes	6	Achievement index, achievement points, recommendation, organizational activity, semester level, and completeness of documents	Scholarship recipient	Yes
Vadim Demichev et al. (2022) [41]	No	No	No	0	-	Optimization of treatment for Covid-19 patients	Yes
Our research	Yes	Yes	Yes	12	Pneumonia, ARDS, CHF, AKI, CAD, Dyspnea, NSTEMI, ADHF, HHD, Febris, Anosmia, Ageusia	Care status of Covid-19 patients, whether inpatient or self-isolation	Yes

Note: ML = Machine Learning

III. METHODOLOGY

This study applies two data mining methods or machine learning algorithms: RF and SVM. The big data is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The big data source used in this study is primary data from patient medical records/documents. The attributes of the patient's disease symptoms and care status classes amount to thousands of patient medical record data. Patient datasets containing non-Covid-19 and duplicate and incomplete Covid-19 patient data are removed, so only data is left as a dataset for data mining processes. Medical record data of disease symptoms obtained

from string data is then converted into numeric data. The development of the application program in this study uses the Python computer programming language.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods. The data mining process in this research uses CRISP-DM (Cross-Industry Standard Process for Data Mining). CRISP-DM is a standard data mining process. The process in CRISP-DM comprises a six-stage [42], as shown in Figure 1 [43]. Figure 2 shows the process carried out at each stage of CRISP-DM.

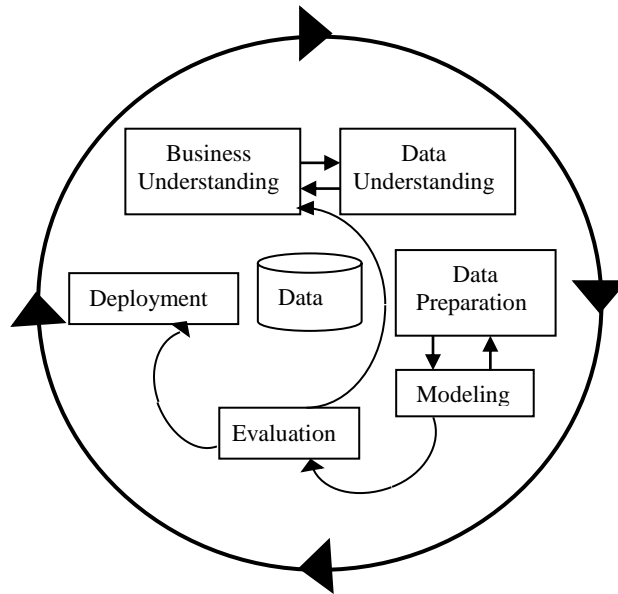


Figure 1. The CRISP-DM Process

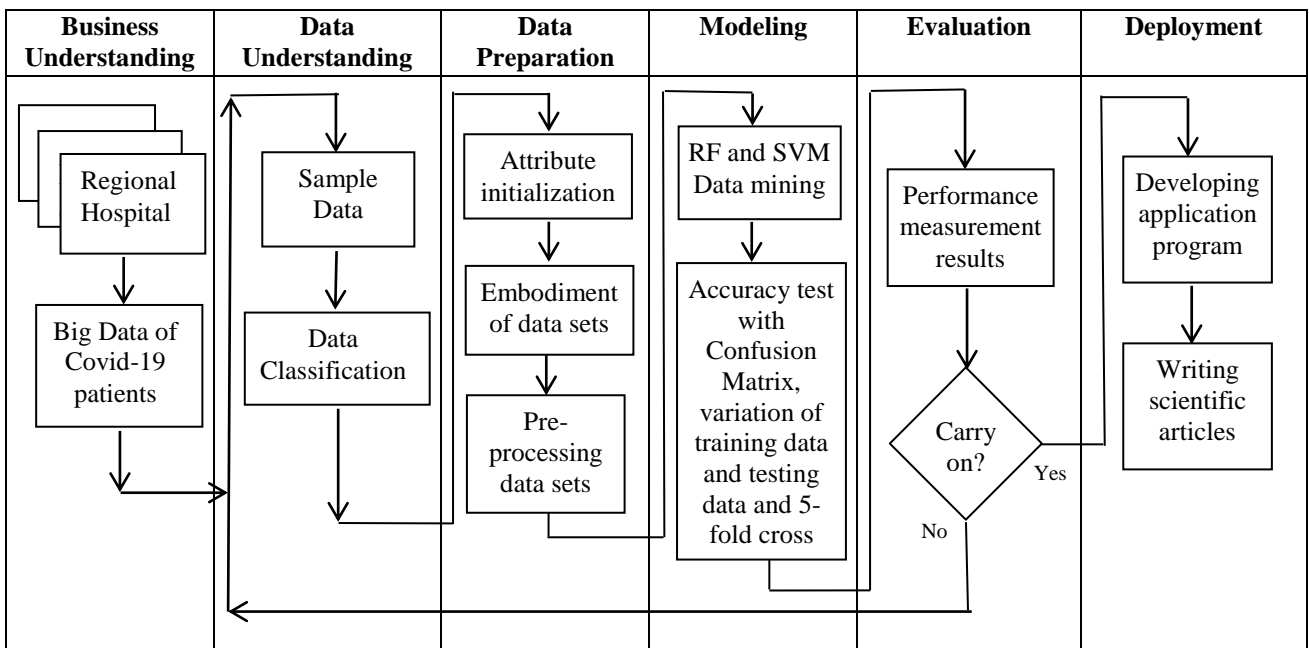


Figure 2. Data Mining Process of Covid-19 Patient Big Data

In Figure 2, business understanding is the stage of sorting out thousands of hospital patient medical data to collect the required patient data. The next stage is understanding the data collected as representative data for Covid-19 patients. This Covid-19 patient data classifies the patient's signs and symptoms and treatment status, which needs further processing at the next stage. The next stage is the data preparation stage, which essentially determines the attributes of the names of signs and symptoms of Covid-19 patients. The embodiment of the dataset containing knowledge according to treatment status refers to the signs and symptoms that the patient has (marked with Yes) or does not have (marked with No). The next thing to do is preprocess the dataset, changing the category value of the symptom attribute and the class attribute with the number 1 and the number -1. The process of extracting raw data obtained in the previous process stage is data that is further processed by data mining methods or used as learning machine learning data at the modeling stage. So that machine learning can predict. The next stage is the evaluation stage, namely, knowing the predictive reliability of the data mining or machine learning method. Then the last stage is the deployment stage to disseminate research results so that they are helpful for implementation by various parties, especially hospitals and other professionals, in the form of developing application programs and scientific articles.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected from medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. There are thousands of data on Covid-19 patients. The patient dataset containing incomplete data and non-covid-19 was omitted or ignored. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 2 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE II. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses. There were 117 patients with Covid-19 who had a different diagnosis from the others. In this study, the number of signs and symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 3).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or G_i where $i = 1, 2, 3 \dots 12$) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 4. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or symptoms.

Furthermore, the preprocessing of the data set is done by changing the G_i with x_i and the G_i Yes attribute value with the number 1 while the G_i No attribute with the number -1. In addition, dataset preprocessing is also carried out on class attributes, namely changing the independent isolation class attribute category with the number 1 and the inpatient class attribute category with the number -1, as shown in Table 5.

TABLE III. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE IV. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	Yes	No	No	No	No	No	Inpatient
2	Yes	Yes	No	Yes	No	No	Inpatient
3	Yes	No	Yes	No	No	No	Inpatient
4	Yes	No	No	No	No	No	Inpatient
5	Yes	No	No	No	No	No	Inpatient
6	Yes	No	No	No	Yes	Yes	Inpatient
7	No	No	Yes	No	No	No	Inpatient
..
..
114	No	No	No	No	No	Yes	Self-isolation
115	Yes	No	No	No	Yes	No	Inpatient
116	Yes	No	No	No	Yes	Yes	Inpatient
117	No	No	No	No	No	No	Inpatient

TABLE V. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
..
..
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [44], which has their respective advantages in building application programs [45][46]. The application program built in this research uses the Python programming language to facilitate patient care status prediction. The hyperparameter used in the SVM method is kernel = linear and Probability = True. Meanwhile, the hyperparameter used for the RF hyperparameter method is n_estimators = 100. The n_estimators is the number of trees in RF.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

In SVM, there are two implementation models: mathematical programming techniques and kernel functions. This study applies kernel functions and focuses on classifying two categories of class attributes. The class attribute is a treatment for $y_i = +1, -1$. The formula of the SVM data mining method is: (a) to form a linear equation from the training data; (b) find the value of w and b, and (c) the value of the classification decision is as follows.

$$S = ((x1, y1), \dots, (xl, yl)) \quad (1)$$

$$y_i ((w \cdot x_i) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

RF uses bootstrap samples from training data to create a tree from a randomly selected subset. The chosen predictor is a candidate for splitting the decision tree. The results of the category predictions from the treatment class based on the results of the highest voting were chosen as the final prediction results. The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_{i=1}^c p_i^2 \quad (4)$$

$$Entropy(S) = \sum_{i=1}^c -p_i \times \log_2(p_i) \quad (5)$$

Description:

S = Set of cases

p_i = the proportion of case i to the Set of cases

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and Recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

Accuracy states the closeness of the measurement results to the actual value, while Precision shows how close the difference in the measurement results is on repeated measurements. On the other hand, Recall states the level of success in retrieving information. Precision and Recall are necessary because Precision denotes a measure of quality, and Recall denotes a measure of quantity.

Measurement of accuracy is based on the ratio between the correct predictions (positive and negative) with the overall data. In contrast, precision measurements are based on the percentage of true positive predictions compared to overall positive predicted outcomes. Meanwhile, the recall measurement is based on the ratio of true positive predictions compared to the general actual positive data.

The format of the confusion matrix table is as shown in Table 6.

TABLE VI. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-Validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation helps assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold value. In the

performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each subgroup to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the training set (Dtrain). In the second fold, the second subset is the validate set, the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 7 and Figure 3. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE VII. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TESTING DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

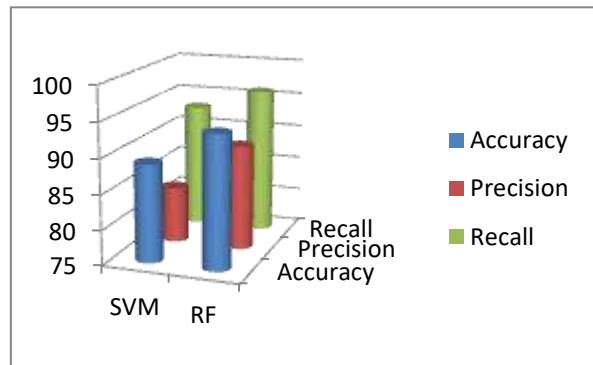


Figure 3. System Model Performance Testing with 85% of Training Data and 15% of Testing Data

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% data 90% testing and training data and 10% testing data are as shown in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)		Accuracy (in %)		Precision (in %)		Recall (in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix. 333

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 9 and Figure 4.

TABLE IX. PREDICTION PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

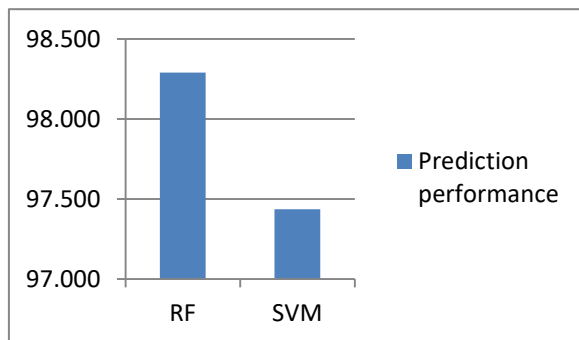


Figure 4. Predictive performance testing using K-fold cross-validation

4.6 Deployment

One of the deployments in this research is making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Thus, the results obtained can be developed and become the knowledge of many parties as a responsibility for the correctness of the effects of research carried out as professional researchers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 90%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%. The research result implication is that RF machine learning can help or replace the role of medical personnel in predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, with high accuracy.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Further research needs to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion for future study is: to conduct further research using other data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients and various other diseases, to build a system that not only predicts but also performs clustering, association, and estimates of various other fields of science, including patients' care status, with a combination of machine learning and the Internet of Things.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first, second, and so on as the fifth author.

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Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

By Anthony Anggrawan

Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

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Abstract—Covid-19 has become a global pandemic that causes many deaths, so medical treatment for Covid-19 patients gets special attention, whether hospitalized or self-isolated. However, the problem in medical action is not easy, and the most frequent mistakes are due to inaccuracies in medical decision-making. Meanwhile, machine learning can predict with high accuracy. For that or that's why this study aims to propose a data mining classification method as a machine learning model to accurately predict the treatment status of Covid-19 patients, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients. The implication is that RF machine learning can help/replace the role of medical experts in predicting the patient's care status.

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Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analysis or better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

In essence, the care status for Covid-19 patients is self-isolation for patients with non-severe illness status and hospitalization for patients who are seriously ill and at critical risk or cause death. Hospitals or medical doctors take various ways to reduce the number of deaths of Covid-19 patients, including by regulating the status of services in hospitals. Expert doctors recommend that patients self-isolate or should be hospitalized after analyzing the patient's medical data. Determining the level of care for Covid-19 patients is a form of medical treatment or treatment for Covid-19 patients to get proper treatment or care.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [19][20][21]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [22]. For this reason (or

why), this study's objective is to propose a machine learning system model for decision-making solutions (predictions) for the treatment status of Covid-19 patients, both inpatient and self-isolation, using data mining methods.

The implication is that the proposed machine learning system model can help and even replace the role of medical experts (specialist doctors) in making medical decisions for Covid-19 patients, whether hospitalization or self-isolation. That is, machine learning performs tasks like a medical specialist in determining the results of the diagnosis of the nutritional status of Covid-19 patients based on medical data of Covid-19 patients. Furthermore, machine learning can work tirelessly, time and place, and has intelligence like an expert, so it is not surprising that previous research confirms that intelligent machines can make superior decisions to experts because humans have a human error factor [23].

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [24]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting new data with high accuracy [22]. After all, predicting individuals with symptoms of being infected with Covid-19 mandates machine learning (application-based) and contributes to effectively isolating Covid-19 patients [25].

Big data demands large storage media [26]. However, big data is no longer traditionally processed [27]. Instead, today's big data processing relies on machines that can provide systematic results [28]. Big data storage is generally on a computer server with a large storage capacity. Still, servers also make it happen by renting online cloud data storage services such as Amazon Simple Storage Service (Amazon S3) and Google Cloud [29]. Cloud facilitates cost-effective big data storage and analysis [29].

Big data processing techniques in data mining include several stages: target data, preprocessed data, data mining, and evaluation/analysis of mining results [29]. Target data and preprocessed data are the processes of extracting raw data from big data [29]. Target data is to select the required data (sample data) and classify data. The preprocessed data is to prepare data sets for data mining, including cleaning up incomplete data, duplicate data, and converting string data into numeric coded data. Finally, data mining and evaluation results extract hidden information by applying data mining methods suitable for the objectives and analyzing them.

Many Data Mining methods include K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative

Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [30] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [31] [32], as well as SVM algorithm has a widely known technique used for classification [32]. It is why this study uses SVM and RF to classify treatment status. Given that the SVM and RF machine learning algorithms are both popularly used by many researchers, the SVM and RF machine learning algorithms are the most appropriate combination used in other research, including research to classify and/or predict the treatment status of Covid-19 patients.

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some related works from the latest scientific articles compared with the work in this research article.

Askin Kavzoglu, İsmail Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms such as RF, SVM, and decision tree classifier (DT) [30]. This previous research is different from the research in the article on the research objectives and the object under study. In the meantime, Celestina Wendi et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [33]. The

2 difference between previous research and the research in this article is that the previous research 4 only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.

Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [34]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, 84 research in this article predicts the treatment status of Covid-19 patients.

Based on patient 19 clinical data, using statistical methods, Boran Hao et al. (2020) developed a model to predict pneumonia severity in Covid-19 patients using the Natural Language Processing tool [35]. However, this previous research differs in the research purpose and way compared to the research in this article. Meanwhile, Anthony Anggrawan et al. 1 (2021) implemented machine learning to diagnose drug users and types of drug-using 11 Forward Chaining and Certainty Factor methods [22]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM. 72

Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson and Gamma distribution [36]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research 19 in this article. In the meantime, Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research object 50 and techniques used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should 4 be hospitalized or self-isolated. This article's research does not use the DT method but uses the RF and SVM methods.

26 Soham Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [37]. This previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not utilizing the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods. Whereas,

Ankit Mehrotra and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [38]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from this article's research because it is an experimental study, not a literature review research.

49 Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods compared to this article's research. The difference 11 is that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM 2 Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data 78 mining methods [14].

Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for 2 scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [39]. However, the previous research differs in the purpose and way compared to this article's research. In contrast, Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of serious 5 ill Covid-19 patients with plasma proteomics [40]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

Table 1 shows a comparison between some of the most recent previous 5 work with the work carried out in this study. By referring to the elaboration of the most recent previous related work by some researchers, the research carried out in this article has novelties (from the last research gap) that previous researchers have not studied. In essence, the gap in earlier research 83 is that no one has researched machine learning models to predict the inpatient status or self-isolation of Covid-19 patients by involving RF and SVM algorithms. In addition, the 12 criteria used to indicate the treatment status of Covid-19 patients are completely different from previous similar studies (as shown in Table 1 in the Criteria/Attributes column). So, the 28 y's originality lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

TABLE I. COMPARISON OF THIS ARTICLE'S WORK WITH SOME PREVIOUS RELATED WORKS

Research by	Research methods			Criteria/Attributes		Research Object	Accuracy Test
	RF	SVM	ML	Number	Name		
Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) [30]	Yes	Yes	Yes	10	58 Coastal Aerosol, Blue, Green, Red, Vegetation Red Edge, NIR, Narrow NIR, Water vapor, SWIR-Cirrus, SWIR	Satellite remote sensing images	Yes
Celestine Iwendi et al. (2020) [33]	Yes	No	Yes	6	Symptom1, symptom2, symptom3, symptom4, symptom5, symptom6	Illness severity of Covid-19 patients	No
Chelvian Aroef, Yuda Rivian, and Zuherman Rustam (2020) [34]	Yes	Yes	Yes	9	76 Age, Body Mass Index (BMI), Glucose, Insulin, Homa, Leptin, Adiponectin, Resistin, MCP 1	Breast cancer	Yes
Boran Hao et al. (2020) [35]	No	No	No	10	Radiology Opacities, Respiratory Rate, Age, Fever Male, Albumin, Anion Gap, SpO2, LDH, Calcium	The severity of pneumonia in Covid-19 patients	Yes
Anthony Anggrawan et al. (2021) [22]	No	No	Yes	27	Out of breath, Anxious, Nausea, Diarrhea, Convulsions, Easily angry, Depression, Sleep patterns change, Eating, Chills, Shaking, Insomnia, Fast heart rate, Blood pressure rises, Difficult to focus, Difficult to rest, Weight loss, Dry mouth, Blurred vision, Changed skin color, Constipation, Stomachache, Drowsiness, Itching, Difficulty urinating, Mood swings, Dizziness	Drug users and types of drug-using	Yes
Hongwei Zhao et al. (2021) [36]	No	No	No	0	-	Number of cases of Covid-19 patients	No
Bassam Mahboub et al. (2021) [8]	No	No	Yes	5	26 Age, Gender, Nationality, Blood group, BMI	The treatment period for Covid-19 patient	Yes
Soham Guhathakurata et al. (2021) [37]	No	Yes	Yes	8	Temp, Breathing rate, Hypertension, Heartbeat rate (HBR), Acute respiratory disease syndrome (ARDS), Chest pain, Heart disease, Cough with sputum (CWS)	Predicting whether patients are infected with Covid-19 or not	No
Ankit Mehrotraa and Reeti Agarwal (2021) [38]	No	No	No	0	-	Discussing the data mining method's role in the Covid-19 pandemic	No
Pratiyush Guleria et al. (2022) [14]	No	Yes	Yes	0	-	The death rate of Covid-19 patient	Yes
Anthony Anggrawan, et al. (2022) [39]	No	No	Yes	6	2 Achievement index, achievement points, recommendation, organizational activity, semester level, and completeness of documents	Scholarship recipient	Yes
Vadim Demichev et al. (2022) [40]	No	No	No	0	-	Optimization of treatment for Covid-19 patients	Yes
Our research	Yes	Yes	Yes	12	Pneumonia, ARDS, CHF, AKI, CAD, Dyspnea, NSTEMI, ADHF, HHD, Febris, Anosmia, Ageusia	Care status of Covid-19 patients, whether inpatient or self-isolation	Yes

Note: ML = Machine Learning

III. METHODOLOGY

This study applies two data mining methods or machine learning algorithms: RF and SVM. The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The big data source used in this study is primary data from patient medical records/documents. The attributes of the patient's disease symptoms and care status classes amount to thousands of patient medical record data. Patient datasets containing non-Covid-19 and duplicate and incomplete Covid-19 patient data are removed, so only data is left as a dataset for data mining processes.

Medical record data of disease symptoms obtained from string data is then converted into numeric data. The development of the application program in this study uses the Python programming language.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods. The methodology in this study uses CRISP-DM (Cross-Industry Standard Process for Data Mining). CRISP-DM is standard data mining. The process in CRISP-DM comprises a six-stage [41], see Figure 1 [42]. Figure 2 shows the data mining processes in this study.

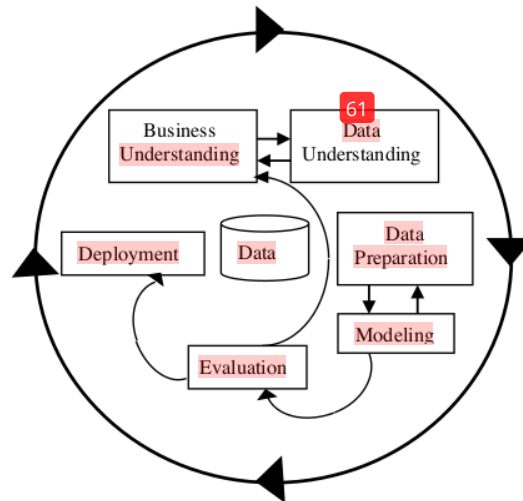


Figure 1. The CRISP-DM Process

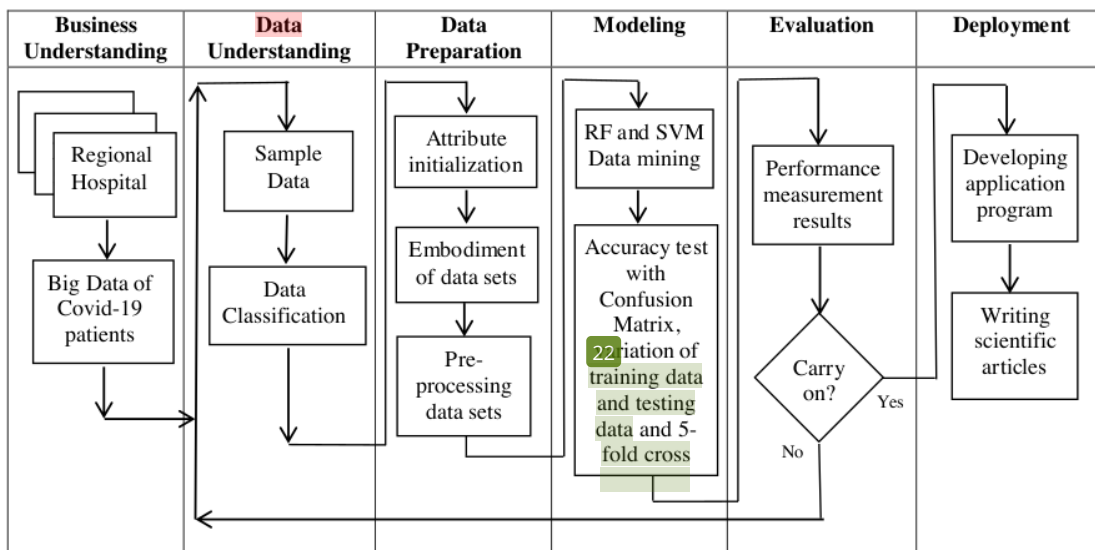


Figure 2. Data Mining Process of Covid-19 Patient Big Data

Business understanding is the stage of sorting out thousands of hospital patient medical data to collect the required patient data. The next stage is understanding the data collected as representative data for Covid-19 patients. This Covid-19 patient data classifies the patient's signs and symptoms and treatment status, which needs further processing at the next stage. The next stage is the data preparation stage, which essentially determines the attributes of the names of signs and symptoms of Covid-19 patients. The embodiment of the dataset containing knowledge according to treatment status refers to the signs and symptoms that the patient has (marked with Yes) or does not have (marked with No). The next thing to do is preprocess the dataset, changing the category value of the symptom attribute and the class attribute with the number 1 and the number -1. The process of extracting raw data obtained in the previous process stage is data that is further processed by data mining methods or used as learning machine learning data at the modeling stage. So that machine learning can predict. The next stage is the evaluation stage, namely, knowing the predictive reliability of the data mining or machine learning method. Then the last stage is the deployment stage to disseminate research results so that they are helpful for implementation by various parties, especially hospitals and other professionals, in the form of developing application programs and scientific articles.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected from medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. There are thousands of data on Covid-19 patients. The patient dataset containing incomplete data and non-covid-19 was omitted/ignored. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

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4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 2 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE II. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses. There were 117 patients with Covid-19 who had a different diagnosis from the others. In this study, the number of signs, symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 3).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or Gi where i = 1, 2,3 ...12) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 4. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or symptoms.

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1. In addition, dataset preprocessing is also carried out on class attributes, namely changing the independent isolation class attribute category with the number 1 and the inpatient class attribute category with the number -1, as shown in Table 5.

TABLE III. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
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G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE IV. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	Yes	No	No	No	No	No	Inpatient
2	Yes	Yes	No	Yes	No	No	Inpatient
3	Yes	No	Yes	No	No	No	Inpatient
4	Yes	No	No	No	No	No	Inpatient
5	Yes	No	No	No	No	No	Inpatient
6	Yes	No	No	No	Yes	Yes	Inpatient
7	No	No	Yes	No	No	No	Inpatient
...
...
114	No	No	No	No	No	Yes	Self-isolation
115	Yes	No	No	No	Yes	No	Inpatient
116	Yes	No	No	No	Yes	Yes	Inpatient
117	No	No	No	No	No	No	Inpatient

TABLE V. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
...
...
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [43], which has

their respective advantages in building application programs [44][45]. The application program built in this research uses the Python programming language to facilitate patient care status prediction. The hyperparameter used in the SVM method is kernel = linear and Probability = True. Meanwhile, the hyperparameter used for the RF hyperparameter method is n_estimators = 100. The n_estimators is the number of trees in RF.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

In SVM, there are two implementation models: mathematical programming techniques and kernel functions. This study applies kernel functions and focuses on classifying two categories of class attributes. The class attribute is a treatment for $y_i = +1, -1$. The formula of the SVM data mining method is: (a) to form a linear equation from the training data; (b) find the value of w and b, and (c) the value of the classification decision is as follows.

$$((x_1, y_1), \dots, (x_l, y_l)) \quad (1)$$

$$y_i ((w \cdot x_i) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S = set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

RF uses bootstrap samples from training data to create a tree from a randomly selected subset. The chosen predictor is a candidate for splitting the decision tree. The results of the category predictions from the treatment class based on the results of the highest voting were chosen as the final prediction results. The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_{i=1}^c p_i^2 \quad (4)$$

$$Entropy(S) = \sum_{i=1}^c -p_i \times \log_2(p_i) \quad (5)$$

Description:

S = Set of cases

p_i = the proportion of case i to the Set of cases

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion

matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and Recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

Measurement of accuracy is based on the ratio between the correct predictions (positive and negative) with the overall data. In contrast, precision measurements are based on the percentage of true positive predictions compared to overall positive predicted outcomes. Meanwhile, the recall measurement is based on the ratio of true positive predictions compared to the general actual positive data.

The format of the confusion matrix table is as shown in Table 6.

TABLE VI. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation helps assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold value. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each subgroup to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the training set (Dtrain). In the second fold, the second subset is the validate set the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 7 and Figure 3. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE 22. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TESTING DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

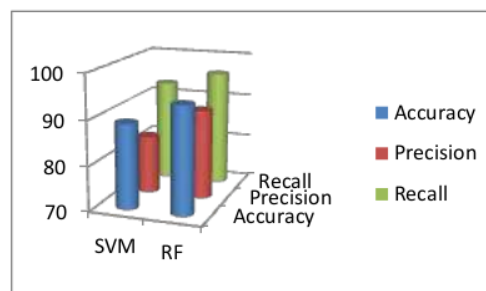


Figure 3. System Model Performance Testing with 85% of Training Data and 15% of Testing Data

Further comparison of the accuracy, precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% testing data and 90% testing and training data and 10% testing data are as shown in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)	Accuracy (in %)	Precision (in %)	Recall (in %)	
Training Testing	RF SVM	RF SVM	RF SVM	
50	50	95 97	97 96	91 96
60	40	96 91	97 90	93 90
70	30	94 92	96 91	92 90
80	20	96 92	94 89	97 94
90	10	92 83	75 67	95 91
Average		95 91	92 87	94 92

Predicting with various test data and training data variations shows that the RF machine learning method

is more accurate and precise than the SVM machine learning method [28] other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix. 333

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 9 and Figure 4.

TABLE IX. PREDICTION PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

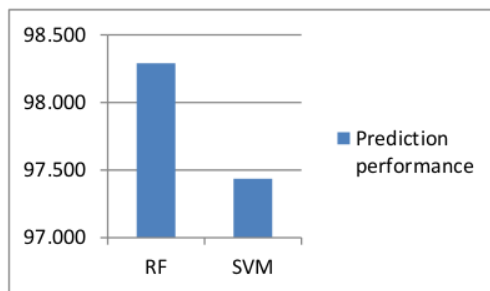


Figure 4. Predictive performance testing using K-fold cross validation

4.6 Deployment

One of the deployments in this research is making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Thus, the results obtained can be developed and become the knowledge of many parties as a responsibility for the correctness of the effects of research carried out as professional researchers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and Precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 90%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction

accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%. The research result implies that RF machine learning can help or replace the role of medical personnel in predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, with high accuracy.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Further research needs to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion for future study is: to conduct further research using other data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients and various other diseases, to build a system that not only predicts but also performs clustering, association, and estimates of various other fields of science, including patients' care status, with a combination of machine learning and the Internet of Things.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first, second, and so on as the fifth author.

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Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

By Anthony Anggrawan

Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

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Abstract—Covid-19 has become a global pandemic that causes many deaths, so medical treatment for Covid-19 patients gets special attention, whether hospitalized or self-isolated. However, the problem in medical action is not easy, and the most frequent mistakes are due to inaccuracies in medical decision-making. Meanwhile, machine learning can predict with high accuracy. For that or that's why this study aims to propose a data mining classification method as a machine learning model to accurately predict the treatment status of Covid-19 patients, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients. The implication is that RF machine learning can help/replace the role of medical experts in predicting the patient's care status.

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Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analysis or better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

In essence, the care status for Covid-19 patients is self-isolation for patients with non-severe illness status and hospitalization for patients who are seriously ill and at critical risk or cause death. Hospitals or medical doctors take various ways to reduce the number of deaths of Covid-19 patients, including by regulating the status of services in hospitals. Expert doctors recommend that patients self-isolate or should be hospitalized after analyzing the patient's medical data. Determining the level of care for Covid-19 patients is a form of medical treatment or treatment for Covid-19 patients to get proper treatment or care.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [19][20][21]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [22]. For this reason (or

why), this study's objective is to propose a machine learning system model for decision-making solutions (predictions) for the treatment status of Covid-19 patients, both inpatient and self-isolation, using data mining methods.

The implication is that the proposed machine learning system model can help and even replace the role of medical experts (specialist doctors) in making medical decisions for Covid-19 patients, whether hospitalization or self-isolation. That is, machine learning performs tasks like a medical specialist in determining the results of the diagnosis of the number and status of Covid-19 patients based on medical data of Covid-19 patients. Furthermore, machine learning can work tirelessly, time and place, and has intelligence like an expert, so it is not surprising that previous research confirms that intelligent machines can make superior decisions to experts because humans have a human error factor [23].

Machine learning can predict classification to predict class membership at regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [24]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting new data with high accuracy [22]. After all, predicting individuals with symptoms of being infected with Covid-19 mandates machine learning (application-based) and contributes to effectively isolating Covid-19 patients [25].

Big data demands large storage media [26]. However, big data is no longer traditionally processed [27]. Instead, today's big data processing relies on machines that can provide systematic results [28]. Big data storage is generally on a computer server with a large storage capacity. Still, scientists also make it happen by renting online cloud data storage services such as Amazon Simple Storage Service (Amazon S3) and Google Cloud [29]. Cloud facilitates cost-effective big data storage and analysis [29].

Big data processing techniques in data mining include several stages: target data, preprocessed data, data mining, and evaluation/analysis of mining results [29]. Target data and preprocessed data are the processes of extracting raw data from big data [29]. Target data is to select the required data (sample data) and classify data. The preprocessed data is to prepare data sets for data mining, including cleaning up incomplete data, duplicate data, and converting string data into numeric coded data. Finally, data mining and evaluation results extract hidden information by applying data mining methods suitable for the objectives and analyzing them.

Many Data Mining methods include K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative

Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [30] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [31] [32], as well as SVM algorithm has a widely known technique used for classification [32]. It is why this study uses SVM and RF to classify treatment status. Given that the SVM and RF machine learning algorithms are both popularly used by many researchers, the SVM and RF machine learning algorithms are the most appropriate combination used in other research, including research to classify and/or predict the treatment status of Covid-19 patients.

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some related works from the latest scientific articles compared with the work in this research article.

Askin Kavzoglu, Fıkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [30]. This previous research is different from the research in the article on the research objectives and the object under study. In the meantime, Celestin Iwendi et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [33]. The

1 difference between previous research and the research in this article is that the previous research 10 only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.

Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [34]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, 48 research in this article predicts the treatment status of Covid-19 patients.

Based on patient 7 clinical data, using statistical methods, Boran Hao et al. (2020) developed a model to predict pneumonia severity in Covid-19 patients using the Natural Language Processing tool [35]. However, this previous research differs in the research purpose and way compared to the research in this article. Meanwhile, Anthony Anggrawan et al. 6 (2021) implemented machine learning to diagnose drug users and types of drug-using Forward Chaining and Certainty Factor methods [22]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.

37 Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson and Gamma distribution [36]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article. In the meantime, Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research object 20 and techniques used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should 10 be hospitalized or self-isolated. This article's research does not use the DT method but uses the RF and SVM methods.

9 Soham Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [37]. This previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not utilizing the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods. Whereas,

Ankit Mehrotra and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [38]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from this article's research because it is an experimental study, not a literature review research.

5 Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods compared to this article's research. The difference 14 is that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, 1 Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].

Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for 1 scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [39]. However, the previous research differs in the purpose and way compared to this article's research. In contrast, Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of serious 4 ill Covid-19 patients with plasma proteomics [40]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

Table 1 shows a comparison between some of the most recent previous 4 related work with the work carried out in this study. By referring to the elaboration of the most recent previous related work by some researchers, the research carried out in this article has novelties (from the last research gap) that previous researchers have not studied. In essence, the gap in earlier research 47 is that no one has researched machine learning models to predict the inpatient status or self-isolation of Covid-19 patients by involving RF and SVM algorithms. In addition, the 12 criteria used to indicate the treatment status of Covid-19 patients are completely different from previous similar studies (as shown in Table 1 in the Criteria/Attributes column 5). So, the study's originality lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

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TABLE I. COMPARISON OF THIS ARTICLE'S WORK WITH SOME PREVIOUS RELATED WORKS

Research by	Research methods			Criteria/Attributes		Research Object	Accuracy Test
	RF	SVM	ML	Number	Name		
Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) [30]	Yes	Yes	Yes	10	Coastal Aerosol, Blue, Green, Red, Vegetation Red Edge, NIR, Narrow NIR, Water vapor, SWIR-Cirrus, SWIR	Satellite remote sensing images	Yes
Celestine Iwendi et al. (2020) [33]	Yes	No	Yes	6	Symptom1, symptom2, symptom3, symptom4, symptom5, symptom6	Illness severity of Covid-19 patients	No
Chelvian Aroef, Yuda Rivian, and Zuherman Rustam (2020) [34]	Yes	Yes	Yes	9	Age, Body Mass Index (BMI), Glucose, Insulin, Homa, Leptin, Adiponectin, Resistin, MCP 1	Breast cancer	Yes
Boran Hao et al. (2020) [35]	No	No	No	10	Radiology Opacities, Respiratory Rate, Age, Fever Male, Albumin, Anion Gap, SpO2, LDH, Calcium	The severity of pneumonia in Covid-19 patients	Yes
Anthony Anggrawan et al. (2021) [22]	No	No	Yes	27	Out of breath, Anxious, Nausea, Diarrhea, Convulsions, Easily angry, Depression, Sleep patterns change, eating, Chills, Shaking, Insomnia, Fast heart rate, Blood pressure rises, Difficult to focus, Difficult to rest, Weight loss, Dry mouth, Blurred vision, Changed skin color, Constipation, Stomachache, Drowsiness, Itching, Difficulty urinating, Mood swings, Dizziness	Drug users and types of drug-using	Yes
Hongwei Zhao et al. (2021) [36]	No	No	No	0	-	Number of cases of Covid-19 patients	No
Bassam Mahboub et al. (2021) [8]	No	No	Yes	5	Age, Gender, Nationality, Blood group, BMI	The treatment period for Covid-19 patient	Yes
Soham Guhathakurata et al. (2021) [37]	No	Yes	Yes	8	Temp, Breathing rate, Hypertension, Heartbeat rate (HBR), Acute respiratory disease syndrome (ARDS), Chest pain, Heart disease, Cough with sputum (CWS)	Predicting whether patients are infected with Covid-19 or not	No
Ankit Mehrotraa and Reeti Agarwal (2021) [38]	No	No	No	0	-	Discussing the data mining method's role in the Covid-19 pandemic	No
Pratiyush Guleria et al. (2022) [14]	No	Yes	Yes	0	-	The death rate of Covid-19 patient	Yes
Anthony Anggrawan, et al. (2022) [39]	No	No	Yes	6	Achievement index, achievement points, recommendation, organizational activity, semester level, and completeness of documents	Scholarship recipient	Yes
Vadim Demichev et al. (2022) [40]	No	No	No	0	-	Optimization of treatment for Covid-19 patients	Yes
Our research	Yes	Yes	Yes	12	Pneumonia, ARDS, CHF, AKI, CAD, Dyspnea, NSTEMI, ADHF, HHD, Febris, Anosmia, Ageusia	Care status of Covid-19 patients, whether inpatient or self-isolation	Yes

Note: ML = Machine Learning

III. METHODOLOGY

This study applies two data mining methods or machine learning algorithms: RF and SVM. The big data used is data on Covid-19 patients from a regional hospital in Mataram, Indonesia. The big data source used in this study is primary data from patient medical records/documents. The attributes of the patient's disease symptoms and care status classes amount to thousands of patient medical record data. Patient datasets containing non-Covid-19 and duplicate and incomplete Covid-19 patient data are removed, so only data is left as a dataset for data mining processes.

Medical record data of disease symptoms obtained from string data is then converted into numeric data. The development of the application program in this study uses the Python programming language.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods. The methodology in this study uses CRISP-DM (Cross-Industry Standard Process for Data Mining). CRISP-DM is standard data mining. The process in CRISP-DM comprises a six-stage [41], see Figure 1 [42]. Figure 2 shows the data mining processes in this study.

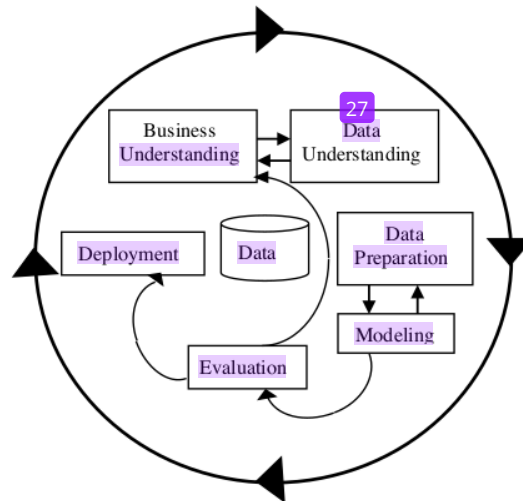


Figure 1. The CRISP-DM Process

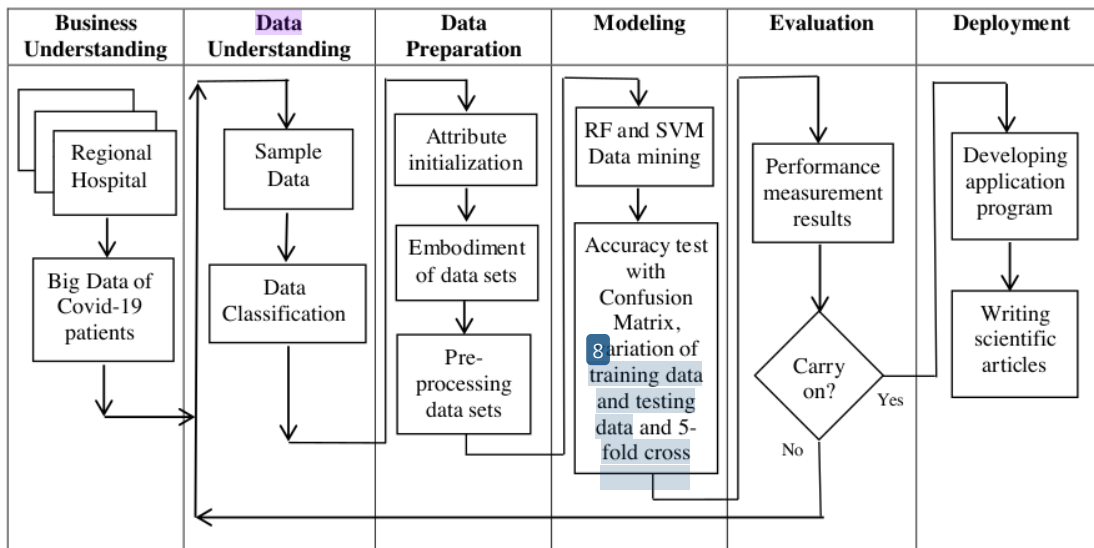


Figure 2. Data Mining Process of Covid-19 Patient Big Data

Business understanding is the stage of sorting out thousands of hospital patient medical data to collect the required patient data. The next stage is understanding the data collected as representative data for Covid-19 patients. This Covid-19 patient data classifies the patient's signs and symptoms and treatment status, which needs further processing at the next stage. The next stage is the data preparation stage, which essentially determines the attributes of the names of signs and symptoms of Covid-19 patients. The embodiment of the dataset containing knowledge according to treatment status refers to the signs and symptoms that the patient has (marked with Yes) or does not have (marked with No). The next thing to do is preprocess the dataset, changing the category value of the symptom attribute and the class attribute with the number 1 and the number -1. The process of extracting raw data obtained in the previous process stage is data that is further processed by data mining methods or used as learning machine learning data at the modeling stage. So that machine learning can predict. The next stage is the evaluation stage, namely, knowing the predictive reliability of the data mining or machine learning method. Then the last stage is the deployment stage to disseminate research results so that they are helpful for implementation by various parties, especially hospitals and other professionals, in the form of developing application programs and scientific articles.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected from medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. There are thousands of data on Covid-19 patients. The patient dataset containing incomplete data and non-covid-19 was omitted/ignored. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

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4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 2 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE II. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses. There were 117 patients with Covid-19 who had a different diagnosis from the others. In this study, the number of signs, symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 3).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or Gi where i = 1, 2,3 ...12) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 4. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or symptoms.

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1. In addition, dataset preprocessing is also carried out on class attributes, namely changing the independent isolation class attribute category with the number 1 and the inpatient class attribute category with the number -1, as shown in Table 5.

TABLE III. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
-----------	-------------------	----------------

G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE IV. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	Yes	No	No	No	No	No	Inpatient
2	Yes	Yes	No	Yes	No	No	Inpatient
3	Yes	No	Yes	No	No	No	Inpatient
4	Yes	No	No	No	No	No	Inpatient
5	Yes	No	No	No	No	No	Inpatient
6	Yes	No	No	No	Yes	Yes	Inpatient
7	No	No	Yes	No	No	No	Inpatient
...
...
114	No	No	No	No	No	Yes	Self-isolation
115	Yes	No	No	No	Yes	No	Inpatient
116	Yes	No	No	No	Yes	Yes	Inpatient
117	No	No	No	No	No	No	Inpatient

TABLE V. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
...
...
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, Known various programming language [43], which has

their respective advantages in building application programs [44][45]. The application program built in this research uses the Python programming language to facilitate patient care status prediction. The hyperparameter used in the SVM method is kernel = linear and Probability = True. Meanwhile, the hyperparameter used for the RF hyperparameter method is n_estimators = 100. The n_estimators is the number of trees in RF.

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

In SVM, there are two implementation models: mathematical programming techniques and kernel functions. This study applies kernel functions and focuses on classifying two categories of class attributes. The class attribute is a treatment for $y_i = +1, -1$. The formula of the SVM data mining method is: (a) to form a linear equation from the training data; (b) find the value of w and b, and (c) the value of the classification decision is as follows.

$$((x_1, y_1), \dots, (x_l, y_l)) \quad (1)$$

$$y_i ((w \cdot x_i) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S = set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

RF uses bootstrap samples from training data to create a tree from a randomly selected subset. The chosen predictor is a candidate for splitting the decision tree. The results of the category predictions from the treatment class based on the results of the highest voting were chosen as the final prediction results. The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_{i=1}^c p_i^2 \quad (4)$$

$$Entropy(S) = \sum_{i=1}^c -p_i \times \log_2(p_i) \quad (5)$$

Description:

S = Set of cases

p_i = the proportion of case i to the Set of cases

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion

matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, precision, and Recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

Measurement of accuracy is based on the ratio between the correct predictions (positive and negative) with the overall data. In contrast, precision measurements are based on the percentage of true positive predictions compared to overall positive predicted outcomes. Meanwhile, the recall measurement is based on the ratio of true positive predictions compared to the general actual positive data.

The format of the confusion matrix table is as shown in Table 6.

TABLE VI. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

4.4.4 K-fold Cross-validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation helps assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold value. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each subgroup to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the training set (Dtrain). In the second fold, the second subset is the validate set the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 7 and Figure 3. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE 8. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TESTING DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

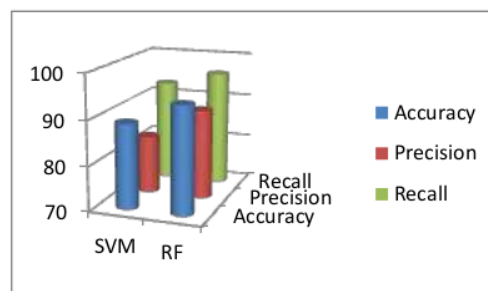


Figure 3. System Model Performance Testing with 85% of Training Data and 15% of Testing Data

Further comparison of the accuracy, precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% testing data and 90% testing and training data and 10% testing data are as shown in Table 8.

TABLE VIII. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)	Accuracy (in %)	Precision (in %)	Recall (in %)				
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method

is more accurate and precise than the SVM machine learning method [5]. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix [44]. 333

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 9 and Figure 4.

TABLE IX. PREDICTION PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

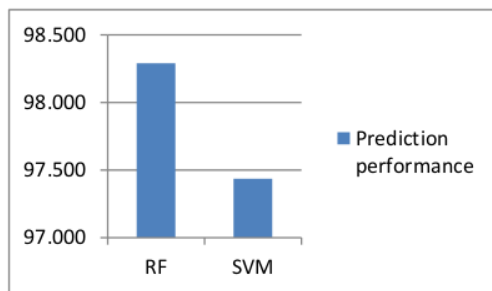


Figure 4. Predictive performance testing using K-fold cross validation

4.6 Deployment

One of the deployments in this research is making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Thus, the results obtained can be developed and become the knowledge of many parties as a responsibility for the correctness of the effects of research carried out as professional researchers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

[11] This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and Precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 90%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction

accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%. The research result implies that RF machine learning can help or replace the role of medical personnel in predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, with high accuracy.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Further research is needed to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion for future study is: to conduct further research using other data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients and various other diseases, to build a system that not only predicts but also performs clustering, association, and estimates of various other fields of science, including patients' care status, with a combination of machine learning and the Internet of Things.

15 CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first, second, and so on as the fifth author.

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We have made improvements to our manuscript following the instructions and suggestions of the reviewers. About which parts we have revised and added writing, it is as explained below:

	Comments to Authors	The Improvements/Corrections (the correction is marked with a yellow highlight)	In-Page																														
1	I suggested the authors to report the hyperparameter details on the ML models. But they are partially addressed. There are many hyperparameters available for SVM and RF (like loss, optimizer, max_iterations, etc.), not just two. This is very crucial for research in ML. I request the authors to report the details in a table.	<p style="text-align: center;">TABLE VI. USE OF HYPERPARAMETER ON SVM AND RF METHOD</p> <table border="1" style="width: 100%; border-collapse: collapse;"> <thead> <tr> <th style="text-align: left;">Classifier Method</th> <th style="text-align: left;">Hyperparameter</th> <th style="text-align: left;">Value</th> </tr> </thead> <tbody> <tr> <td rowspan="7" style="vertical-align: middle;">SVM</td> <td>C</td> <td>1</td> </tr> <tr> <td>Kernel</td> <td>Rbf</td> </tr> <tr> <td>Degree</td> <td>3</td> </tr> <tr> <td>Gamma</td> <td>Scale</td> </tr> <tr> <td>Coef</td> <td>0</td> </tr> <tr> <td>Tol</td> <td>0,000</td> </tr> <tr> <td>Max_iter</td> <td>1</td> </tr> <tr> <td rowspan="5" style="vertical-align: middle;">RF</td> <td>n_estimators</td> <td>100</td> </tr> <tr> <td>Criterion</td> <td>Gini</td> </tr> <tr> <td>Max_depth</td> <td>None</td> </tr> <tr> <td>Min_samples_split</td> <td>2</td> </tr> <tr> <td>Min_samples_leaf</td> <td>1</td> </tr> </tbody> </table>	Classifier Method	Hyperparameter	Value	SVM	C	1	Kernel	Rbf	Degree	3	Gamma	Scale	Coef	0	Tol	0,000	Max_iter	1	RF	n_estimators	100	Criterion	Gini	Max_depth	None	Min_samples_split	2	Min_samples_leaf	1	In-Page 7, 2 nd column	
Classifier Method	Hyperparameter	Value																															
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3	Still some language errors are prevalent	The English language of the manuscript has undergone extensive English editing																															

Hopefully, what we have done fulfills the wishes of the reviewers. Thank you very much.

Sincerely yours
Authors

Plagiarism Check _ Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

By Anthony Anggrawan

Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

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Abstract—Covid-19 has become a global pandemic that causes many deaths, so medical treatment for Covid-19 patients gets special attention, whether hospitalized or self-isolated. However, the problem in medical action is not easy, and the most frequent mistakes are due to inaccuracies in medical decision-making. Meanwhile, machine learning can predict with high accuracy. For that, or that's why this study aims to propose a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients. The implication is that RF machine learning can help/replace the role of medical experts in predicting the patient's care status.

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Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not only because of high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analysis or better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

In essence, the care status for Covid-19 patients is self-isolation for patients with non-severe illness status and hospitalization for patients who are seriously ill and at critical risk or cause death. Hospitals or medical doctors take various ways to reduce the number of deaths of Covid-19 patients, including by regulating the status of services in hospitals. Expert doctors recommend that patients self-isolate or should be hospitalized after analyzing the patient's medical data. Determining the level of care for Covid-19 patients is a form of medical treatment or treatment for Covid-19 patients to get proper treatment or care.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both in-hospital and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18][19], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [20][21][22]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [23]. For this reason (or

why), this study's objective is to propose a machine learning system model for decision-making solutions (predictions) for the treatment status of Covid-19 patients, both inpatient and self-isolation, using data mining methods.

The implication is that the proposed machine learning system model can help and even replace the role of medical experts (specialist doctors) in making medical decisions for Covid-19 patients, whether hospitalization or self-isolation. Machine learning performs tasks like a medical specialist in determining the results of the diagnosis of the treatment status of Covid-19 patients based on medical data of Covid-19 patients. Furthermore, machine learning can work tirelessly, time and place, and has intelligence like an expert, so it is not surprising that previous research confirms that intelligent machines can make superior decisions to experts because humans have a human error factor [24].

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [25]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting new data with high accuracy [23]. After all, predicting individuals with symptoms of being infected with Covid-19 mandates machine learning (application-based) and contributes to effectively isolating Covid-19 patients [26].

Big data demands large storage media [27]. However, big data is no longer traditionally processed [28]. Instead, today's big data processing relies on machines that can provide systematic results [29]. Big data storage is generally on a computer server with a large storage capacity. Still, scientists also make it happen by renting online cloud data storage services such as Amazon Simple Storage Service (Amazon S3) and Google Cloud [30]. Cloud facilitates cost-effective big data storage and analysis [30].

Big data processing techniques in data mining include several stages: target data, preprocessed data, data mining, and evaluation/analysis of mining results [30]. Target data and preprocessed data are the processes of extracting raw data from big data [30]. Target data is to select the required data (sample data) and classify data. The preprocessed data is to prepare data sets for data mining, including cleaning up incomplete data, duplicate data, and converting string data into numeric coded data. Finally, data mining and evaluation results extract hidden information by applying data mining methods suitable for the objectives and analyzing them.

Many Data Mining methods include K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative

Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [31] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [32] [33], as well as SVM algorithm has a widely known technique used for classification [33]. It is why this study uses SVM and RF to classify treatment status. Given that the SVM and RF machine learning algorithms are both popularly used by many researchers, the SVM and RF machine learning algorithms are the most appropriate combination used in research, including research to classify and predict the treatment status of Covid-19 patients.

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some related works from the latest scientific articles compared with the work in this research article.

Askin Kavzoglu, İrfan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [31]. This previous research is different from the research in the article on the research objectives and the object under study. In the meantime, Celestini et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [34]. The

2 difference between previous research and the research in this article is that the previous research 5 only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.

Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [35]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, 81 research in this article predicts the treatment status of Covid-19 patients.

Based on patient 21 clinical data, using statistical methods, Boran Hao et al. (2020) developed a model to predict pneumonia severity in Covid-19 patients using the Natural Language Processing tool [36]. However, this previous research differs in the research purpose and way compared to the research in this article. Meanwhile, Anthony Anggrawan et al. 41 (2021) implemented machine learning to diagnose drug users and types of drug-using Forward Chaining and Certainty Factor methods [23]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM. 68

Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson and Gamma distribution [37]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research 21 in this article. In the meantime, Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research object 26 and techniques used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should 5 be hospitalized or self-isolated. This article's research does not use the DT method but uses the RF and SVM methods.

27 Soham Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [38]. However, this previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not utilizing the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods. At the

same time, Ankit Mehrotra and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [39]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from this article's research because it is an experimental study, not a literature review.

72 Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods compared to this article's research. The difference is 11 at previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM 2, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14]. 75

Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for 2 scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [40]. However, the previous research differs in the purpose and way compared to this article's research. In contrast, Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of serious 3 ill Covid-19 patients with plasma proteomics [41]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

Table 1 compares some of the most recent previous 3 related work with the work carried out in this study. By referring to the elaboration of the most recent last related work by some researchers, the research carried out in this article has novelties (from the prior research gap) that previous researchers have not studied. In essence, the gap in earlier research 11 is that no one has researched machine learning models to predict the inpatient status or self-isolation of Covid-19 patients by involving RF and SVM algorithms. In addition, the 12 criteria used to indicate the treatment status of Covid-19 patients are entirely different from previous similar studies (as shown in Table 1 in the Criteria/Attributes column). So, the 15 study's originality lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

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TABLE I. COMPARISON OF THIS ARTICLE'S WORK WITH SOME PREVIOUS RELATED WORKS

Research by	Research methods			Criteria/Attributes		Research Object	Accuracy Test
	RF	SVM	ML	Number	Name		
Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) [31]	Yes	Yes	Yes	10	Coastal Aerosol, Blue, Green, Red, Vegetation Red Edge, NIR, Narrow NIR, Water vapor, SWIR-Cirrus, SWIR	Satellite remote sensing images	Yes
Celestine Iwendi et al. (2020) [34]	Yes	No	Yes	6	Symptom1, symptom2, symptom3, symptom4, symptom5, symptom6	Illness severity of Covid-19 patients	No
Chelvian Aroef, Yuda Rivin, and Zuherman Rustam (2020) [35]	Yes	Yes	Yes	9	Age, Body Mass Index (BMI), Glucose, Insulin, Homa, Leptin, Adiponectin, Resistin, MCP 1	Breast cancer	Yes
Boran Hao et al. (2020) [36]	No	No	No	10	Radiology Opacities, Respiratory Rate, Age, Fever Male, Albumin, Anion Gap, SpO2, LDH, Calcium	The severity of pneumonia in Covid-19 patients	Yes
Anthony Anggrawan et al. (2021) [23]	No	No	Yes	27	Out of breath, Anxious, Nausea, Diarrhea, Convulsions, Easily angry, Depression, Sleep patterns change, Sweating, Chills, Shaking, Insomnia, Fast heart rate, Blood pressure rises, Difficult to focus, Difficult to rest, Weight loss, Dry mouth, Blurred vision, Changed skin color, Constipation, Stomachache, Drowsiness, Itching, Difficulty urinating, Mood swings, Dizziness	Drug users and types of drug-using	Yes
Hongwei Zhao et al. (2021) [37]	No	No	No	0	-	Number of cases of Covid-19 patients	No
Bassam Mahboub et al. (2021) [8]	No	No	Yes	5	Age, Gender, Nationality, Blood group, BMI	The treatment period for Covid-19 patient	Yes
Soham Guhathakurata et al. (2021) [38]	No	Yes	Yes	8	Temp, Breathing rate, Hypertension, Heartbeat rate (HBR), Acute respiratory disease syndrome (ARDS), Chest pain, Heart disease, Cough with sputum (CWS)	Predicting whether patients are infected with Covid-19 or not	No
Ankit Mehrotraa and Reeti Agarwal (2021) [39]	No	No	No	0	-	Discussing the data mining method's role in the Covid-19 pandemic	No
Pratiyush Guleria et al. (2022) [14]	No	Yes	Yes	0	-	The death rate of Covid-19 patient	Yes
Anthony Anggrawan, et al. (2022) [40]	No	No	Yes	6	Achievement index, achievement points, recommendation, organizational activity, semester level, and completeness of documents	Scholarship recipient	Yes
Vadim Demichev et al. (2022) [41]	No	No	No	0	-	Optimization of treatment for Covid-19 patients	Yes
Our research	Yes	Yes	Yes	12	Pneumonia, ARDS, CHF, AKI, CAD, Dyspnea, NSTEMI, ADHF, HHD, Febris, Anosmia, Ageusia	Care status of Covid-19 patients, whether inpatient or self-isolation	Yes

Note: ML = Machine Learning

III. METHODOLOGY

This study applies two data mining methods or machine learning algorithms: RF and SVM. The big data is on Covid-19 patients from a regional hospital in Mataram, Indonesia. The significant data source used in this study is primary data from patient medical records/documents. The attributes of the patient's disease symptoms and care status classes amount to thousands of patient medical record data. Patient datasets containing non-Covid-19 and duplicate and incomplete Covid-19 patient data are removed, so only data is left as a dataset for data mining processes. Medical record data of disease symptoms obtained

from string data is then converted into numeric data. The development of the application program in this study uses the Python computer programming language.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods. The data mining process in this research uses CRISP-DM (Cross-Industry Standard Process for Data Mining). CRISP-DM is a standard data mining process. The process in CRISP-DM comprises a six-stage [42], as shown in Figure 1 [43]. Figure 2 shows the process carried out at each stage of CRISP-DM.

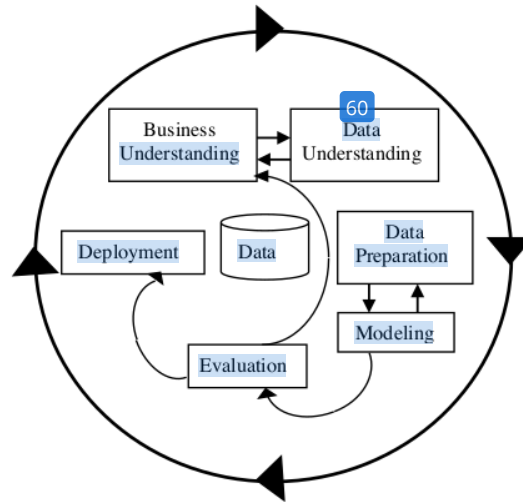


Figure 1. The CRISP-DM Process

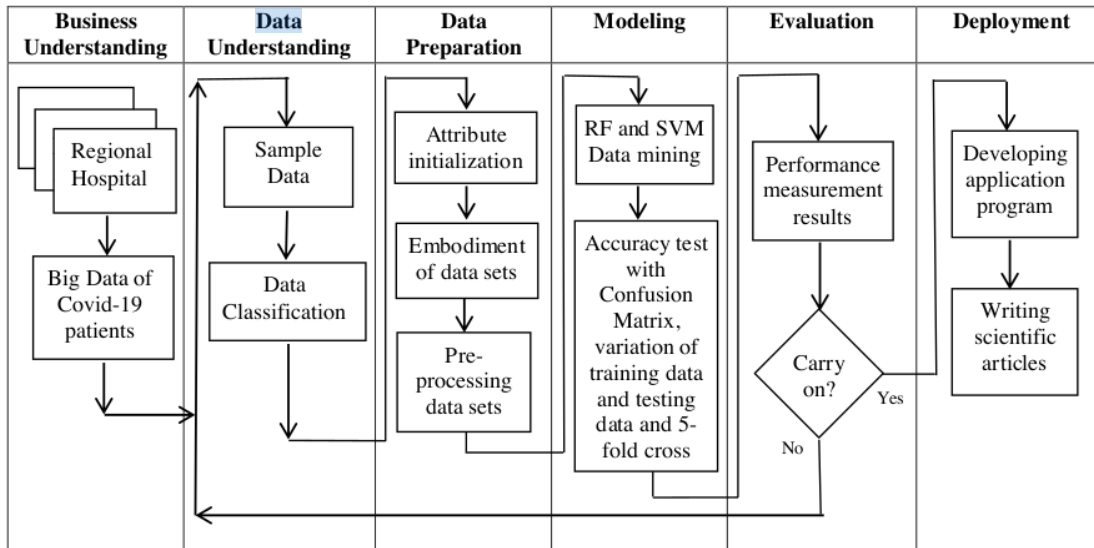


Figure 2. Data Mining Process of Covid-19 Patient Big Data

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 2 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE II. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses. There were 117 patients with Covid-19 who had a different diagnosis from the others. In this study, the number of signs, symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 3).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or G_i where $i = 1, 2, 3 \dots 12$) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 4. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or symptoms.

Furthermore, the preprocessing of the data set is done by changing the G_i with x_i and the G_i Yes attribute value with the number 1 while the G_i No attribute with the number -1. In addition, dataset preprocessing is also carried out on class attributes, namely changing the independent isolation class attribute category with the number 1 and the inpatient class attribute category with the number -1, as shown in Table 5.

In Figure 2, business understanding is the stage of sorting out thousands of hospital patient medical data to collect the required patient data. The next stage is understanding the data collected as representative data for Covid-19 patients. This Covid-19 patient data classifies the patient's signs and symptoms and treatment status, which needs further processing at the next stage. The next stage is the data preparation stage, which essentially determines the attributes of the names of signs and symptoms of Covid-19 patients. The embodiment of the dataset containing knowledge according to treatment status refers to the signs and symptoms that the patient has (marked with Yes) or does not have (marked with No). The next thing to do is preprocess the dataset, changing the category value of the symptom attribute and the class attribute with the number 1 and the number -1. The process of extracting raw data obtained in the previous process stage is data that is further processed by data mining methods or used as learning machine learning data at the modeling stage. So that machine learning can predict. The next stage is the evaluation stage, namely, knowing the predictive reliability of the data mining or machine learning method. Then the last stage is the deployment stage to disseminate research results so that they are helpful for implementation by various parties, especially hospitals and other professionals, in the form of developing application programs and scientific articles.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected from medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. There are thousands of data on Covid-19 patients. The patient dataset containing incomplete data and non-covid-19 was omitted or ignored. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

TABLE III. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE IV. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	Yes	No	No	No	No	No	Inpatient
2	Yes	Yes	No	Yes	No	No	Inpatient
3	Yes	No	Yes	No	No	No	Inpatient
4	Yes	No	No	No	No	No	Inpatient
5	Yes	No	No	No	No	No	Inpatient
6	Yes	No	No	No	Yes	Yes	Inpatient
7	No	No	Yes	No	No	No	Inpatient
...
...
114	No	No	No	No	No	Yes	Self-isolation
115	Yes	No	No	No	Yes	No	Inpatient
116	Yes	No	No	No	Yes	Yes	Inpatient
117	No	No	No	No	No	No	Inpatient

TABLE V. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
...
...
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

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The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, known various programming language [44], which has their respective advantages in building application programs [45][46]. The application program built in this research uses the Python programming language to facilitate patient care status prediction (as show in Table 6).

TABLE VI. USE OF HYPERPARAMETER ON SVM AND RF METHOD

Classifier	Hyperparameter	Value	
SVM	Method		
	Kernel	Rbf	
	Degree	3	
	Gamma	Scale	
	Coef	0	
	Tol	0,000	
	Max_iter	1	
	Max_iter	-1	
	Random Forest	n_estimators	100
	Random Forest	Criterion	Gini
Random Forest	Max_depth	None	
Random Forest	Min_samples_split	2	
Random Forest	Min_samples_leaf	1	

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows [85]: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

In SVM, there are two implementation models: mathematical programming techniques and kernel functions. This study applies kernel functions and focuses on classifying two categories of class attributes. The class attribute is a treatment for $y_i = +1, -1$. The formula of the SVM data mining method is: (a) to form a linear equation from the training data; (b) find the value of w and b, and (c) the value of the classification decision is as follows.

$$((x_1, y_1), \dots, (x_l, y_l)) \quad (1)$$

$$y_i ((w \cdot x_i) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

RF uses bootstrap samples from training data to create a tree from a randomly selected subset. The chosen predictor is a candidate for splitting the decision tree.

The results of the category predictions from the treatment class based on the results of the highest voting were chosen as the final prediction results. The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_{i=1}^c p_i^2 \quad (4)$$

$$Entropy(S) = \sum_{i=1}^c -p_i \times \log_2(p_i) \quad (5)$$

Description:

S = Set of cases

p_i = the proportion of case i to the Set of cases

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4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and Recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

Accuracy states the closeness of the measurement results to the actual value, while Precision shows how close the difference in the measurement results is on repeated measurements. On the other hand, Recall states the level of success in retrieving information. Precision and Recall are necessary because Precision denotes a measure of quality, and Recall denotes a measure of quantity.

Measurement of accuracy is based on the ratio between the correct predictions (positive and negative) with the overall data. In contrast, precision measurements are based on the percentage of true positive predictions compared to overall positive predicted outcomes. Meanwhile, the recall measurement is based on the ratio of true positive predictions compared to the general actual positive data.

The format of the confusion matrix table is as shown in Table 7. The results of the predictions of the SVM

and RF methods are shown in Tables 8 and 9.

TABLE VII. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

TABLE VIII. CONFUSION MATRIX OF SVM

		Prediction	
		Self-isolation	Inpatient
Actual	Class		
	Self-isolation	4	0
Inpatient		2	12

TABLE IX. CONFUSION MATRIX OF RF

		Prediction	
		Self-isolation	Inpatient
Actual	Class		
	Self-isolation	4	0
Inpatient		1	13

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4.4.4 K-fold Cross-Validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation helps assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k -fold value. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets ($k = 5$). It allows each subgroup to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the training set (Dtrain). In the second fold, the second subset is the validate set, the remaining subset is the training set, and so on until the 5th fold.

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4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 8 and Figure 3. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE X. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TESTING DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

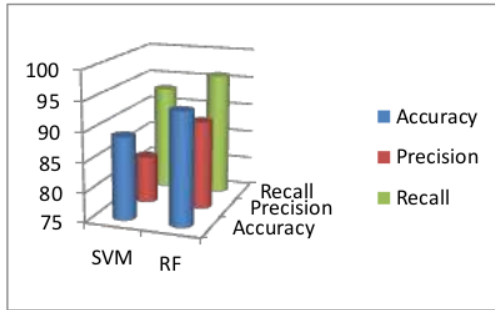


Figure 3. System Model Performance Testing with 85% of Training Data and 15% of Testing Data

Further comparison of the accuracy, precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% testing data and 90% training data and 10% testing data are as shown in Table 11.

TABLE XI. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)		Accuracy (in %)		Precision (in %)		Recall (in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 12 and Figure 4.

TABLE XII. PREDICTION PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

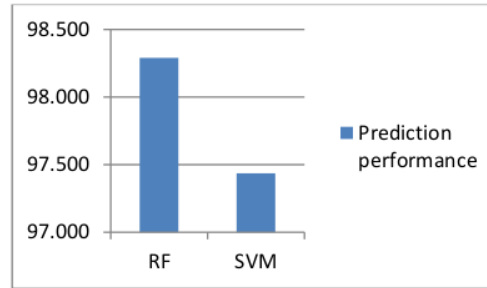


Figure 4. Predictive performance testing using K-fold cross-validation

4.6 Deployment

One of the deployments in this research is making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Thus, the results obtained can be developed and become the knowledge of many parties as a responsibility for the correctness of the effects of research carried out as professional researchers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 90%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%. The research result implies that RF machine learning can help or replace the role of medical personnel in predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, with high accuracy.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Further research needs to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion for future study is: to conduct further research using other data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients and various other diseases, to build a system that not only predicts but also performs clustering, association, and estimates of various other fields of science, including patients' care status, with a combination of

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first, second, and so on as the fifth author.

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Plagiarism Check _ Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

By Anthony Anggrawan

Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

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61

Abstract—Covid-19 has become a global pandemic that causes many deaths, so medical treatment for Covid-19 patients gets special attention, whether hospitalized or self-isolated. However, the problem in medical action is not easy, and the most frequent mistakes are due to inaccuracies in medical decision-making. Meanwhile, machine learning can predict with high accuracy. For that, or that's why this study aims to propose a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients. The implication is that RF machine learning can help/replace the role of medical experts in predicting the patient's care status.

22

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not only because of high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analysis or better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

In essence, the care status for Covid-19 patients is self-isolation for patients with non-severe illness status and hospitalization for patients who are seriously ill and at critical risk or cause death. Hospitals or medical doctors take various ways to reduce the number of deaths of Covid-19 patients, including by regulating the status of services in hospitals. Expert doctors recommend that patients self-isolate or should be hospitalized after analyzing the patient's medical data. Determining the level of care for Covid-19 patients is a form of medical treatment or treatment for Covid-19 patients to get proper treatment or care.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18][19], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [20][21][22]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [23]. For this reason (or

why), this study's objective is to propose a machine learning system model for decision-making solutions (predictions) for the treatment status of Covid-19 patients, both inpatient and self-isolation, using data mining methods.

The implication is that the proposed machine learning system model can help and even replace the role of medical experts (specialist doctors) in making medical decisions for Covid-19 patients, whether hospitalization or self-isolation. Machine learning performs tasks like a medical specialist in determining the results of the diagnosis of the treatment status of Covid-19 patients based on medical data of Covid-19 patients. Furthermore, machine learning can work tirelessly, time and place, and has intelligence like an expert, so it is not surprising that previous research confirms that intelligent machines can make superior decisions to experts because humans have a human error factor [24].

Machine learning can predict classification to predict class membership at regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [25]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting new data with high accuracy [23]. After all, predicting individuals with symptoms of being infected with Covid-19 mandates machine learning (application-based) and contributes to effectively isolating Covid-19 patients [26].

Big data demands large storage media [27]. However, big data is no longer traditionally processed [28]. Instead, today's big data processing relies on machines that can provide systematic results [29]. Big data storage is generally on a computer server with a large storage capacity. Still, services also make it happen by renting online cloud data storage services such as Amazon Simple Storage Service (Amazon S3) and Google Cloud [30]. Cloud facilitates cost-effective big data storage and analysis [30].

Big data processing techniques in data mining include several stages: target data, preprocessed data, data mining, and evaluation/analysis of mining results [30]. Target data and preprocessed data are the processes of extracting raw data from big data [30]. Target data is to select the required data (sample data) and classify data. The preprocessed data is to prepare data sets for data mining, including cleaning up incomplete data, duplicate data, and converting string data into numeric coded data. Finally, data mining and evaluation results extract hidden information by applying data mining methods suitable for the objectives and analyzing them.

Many Data Mining methods include K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative

Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [31] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [32] [33], as well as SVM algorithm has a widely known technique used for classification [33]. It is why this study uses SVM and RF to classify treatment status. Given that the SVM and RF machine learning algorithms are both popularly used by many researchers, the SVM and RF machine learning algorithms are the most appropriate combination used in research, including research to classify and predict the treatment status of Covid-19 patients.

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some related works from the latest scientific articles compared with the work in this research article.

Askin Kavzoglu, İbrahim Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [31]. This previous research is different from the research in the article on the research objectives and the object under study. In the meantime, Celestin Iwendi et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [34]. The

1 difference between previous research and the research in this article is that the previous research 9 only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.

Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [35]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, 47 research in this article predicts the treatment status of Covid-19 patients.

Based on patient 7 clinical data, using statistical methods, Boran Hao et al. (2020) developed a model to predict pneumonia severity in Covid-19 patients using the Natural Language Processing tool [36]. However, this previous research differs in the research purpose and way compared to the research in this article. Meanwhile, Anthony Anggrawan et al. 6 (2021) implemented machine learning to diagnose drug users and types of drug-using 6 Forward Chaining and Certainty Factor methods [23]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.

34 Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson and Gamma distribution [37]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article. In the meantime, Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research object 21 and techniques used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should 9 be hospitalized or self-isolated. This article's research does not use the DT method but uses the RF and SVM methods.

8 Soham Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [38]. However, this previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not utilizing the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods. At the

same time, Ankit Mehrotra and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [39]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from this article's research because it is an experimental study, not a literature review.

38 Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods compared to this article's research. The difference is that previous study 2 examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM 1 Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].

Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for 1 scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [40]. However, the previous research differs in the purpose and way compared to this article's research. In contrast, Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of serious 5 ill Covid-19 patients with plasma proteomics [41]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

Table 1 compares some of the most recent previous 5 related work with the work carried out in this study. By referring to the elaboration of the most recent last related work by some researchers, the research carried out in this article has novelties (from the prior research gap) that previous researchers have not studied. In essence, the gap in earlier research 20 is that no one has researched machine learning models to predict the inpatient status or self-isolation of Covid-19 patients by involving RF and SVM algorithms. In addition, the 12 criteria used to indicate the treatment status of Covid-19 patients are entirely different from previous similar studies (as shown in Table 1 in the Criteria/Attributes 2 column). So, the study's originality lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

1
TABLE I. COMPARISON OF THIS ARTICLE'S WORK WITH SOME PREVIOUS RELATED WORKS

Research by	Research methods			Criteria/Attributes		Research Object	Accuracy Test
	RF	SVM	ML	Number	Name		
Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) [31]	Yes	Yes	Yes	10	Coastal Aerosol, Blue, Green, Red, Vegetation Red Edge, NIR, Narrow NIR, Water vapor, SWIR-Cirrus, SWIR	Satellite remote sensing images	Yes
Celestine Iwendi et al. (2020) [34]	Yes	No	Yes	6	Symptom1, symptom2, symptom3, symptom4, symptom5, symptom6	Illness severity of Covid-19 patients	No
Chelvian Aroef, Yuda Rivin, and Zuherman Rustam (2020) [35]	Yes	Yes	Yes	9	Age, Body Mass Index (BMI), Glucose, Insulin, Homa, Leptin, Adiponectin, Resistin, MCP 1	Breast cancer	Yes
Boran Hao et al. (2020) [36]	No	No	No	10	Radiology Opacities, Respiratory Rate, Age, Fever Male, Albumin, Anion Gap, SpO2, LDH, Calcium	The severity of pneumonia in Covid-19 patients	Yes
Anthony Anggrawan et al. (2021) [23]	No	No	Yes	27	Out of breath, Anxious, Nausea, Diarrhea, Convulsions, Easily angry, Depression, Sleep patterns change, eating, Chills, Shaking, Insomnia, Fast heart rate, Blood pressure rises, Difficult to focus, Difficult to rest, Weight loss, Dry mouth, Blurred vision, Changed skin color, Constipation, Stomachache, Drowsiness, Itching, Difficulty urinating, Mood swings, Dizziness	Drug users and types of drug-using	Yes
Hongwei Zhao et al. (2021) [37]	No	No	No	0	-	Number of cases of Covid-19 patients	No
Bassam Mahboub et al. (2021) [8]	No	No	Yes	5	Age, Gender, Nationality, Blood group, BMI	The treatment period for Covid-19 patient	Yes
Soham Guhathakurata et al. (2021) [38]	No	Yes	Yes	8	Temp, Breathing rate, Hypertension, Heartbeat rate (HBR), Acute respiratory disease syndrome (ARDS), Chest pain, Heart disease, Cough with sputum (CWS)	Predicting whether patients are infected with Covid-19 or not	No
Ankit Mehrotraa and Reeti Agarwal (2021) [39]	No	No	No	0	-	Discussing the data mining method's role in the Covid-19 pandemic	No
Pratiyush Guleria et al. (2022) [14]	No	Yes	Yes	0	-	The death rate of Covid-19 patient	Yes
Anthony Anggrawan, et al. (2022) [40]	No	No	Yes	6	Achievement index, achievement points, recommendation, organizational activity, semester level, and completeness of documents	Scholarship recipient	Yes
Vadim Demichev et al. (2022) [41]	No	No	No	0	-	Optimization of treatment for Covid-19 patients	Yes
Our research	Yes	Yes	Yes	12	Pneumonia, ARDS, CHF, AKI, CAD, Dyspnea, NSTEMI, ADHF, HHD, Febris, Anosmia, Ageusia	Care status of Covid-19 patients, whether inpatient or self-isolation	Yes

Note: ML = Machine Learning

III. METHODOLOGY

This study applies two data mining methods or machine learning algorithms: RF and SVM. The big data is on Covid-19 patients from a regional hospital in Mataram, Indonesia. The significant data source used in this study is primary data from patient medical records/documents. The attributes of the patient's disease symptoms and care status classes amount to thousands of patient medical record data. Patient datasets containing non-Covid-19 and duplicate and incomplete Covid-19 patient data are removed, so only data is left as a dataset for data mining processes. Medical record data of disease symptoms obtained

from string data is then converted into numeric data. The development of the application program in this study uses the Python computer programming language.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods. The data mining process in this research uses CRISP-DM (Cross-Industry Standard Process for Data Mining). CRISP-DM is a standard data mining process. The process in CRISP-DM comprises a six-stage [42], as shown in Figure 1 [43]. Figure 2 shows the process carried out at each stage of CRISP-DM.

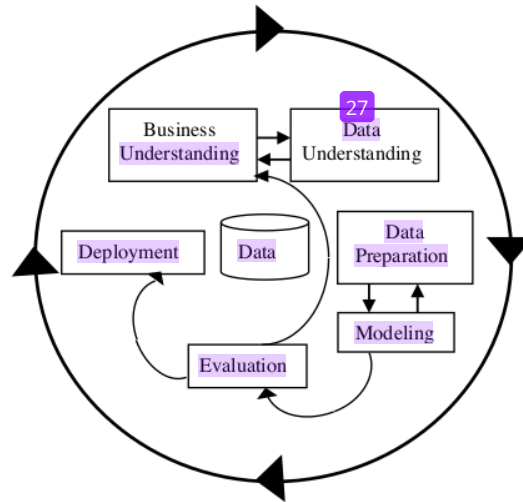


Figure 1. The CRISP-DM Process

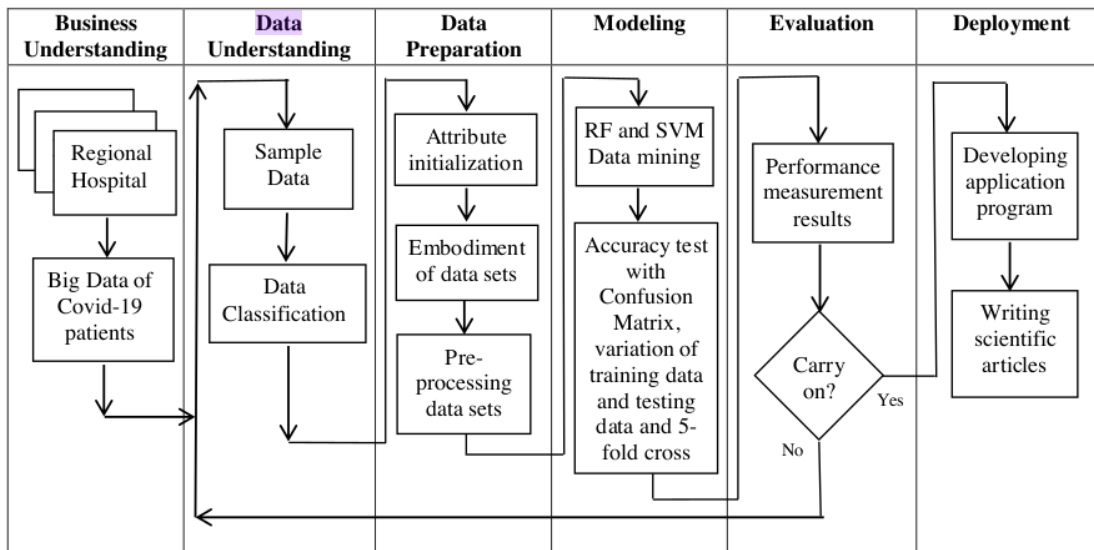


Figure 2. Data Mining Process of Covid-19 Patient Big Data

In Figure 2, business understanding is the stage of sorting out thousands of hospital patient medical data to collect the required patient data. The next stage is understanding the data collected as representative data for Covid-19 patients. This Covid-19 patient data classifies the patient's signs and symptoms and treatment status, which needs further processing at the next stage. The next stage is the data preparation stage, which essentially determines the attributes of the names of signs and symptoms of Covid-19 patients. The embodiment of the dataset containing knowledge according to treatment status refers to the signs and symptoms that the patient has (marked with Yes) or does not have (marked with No). The next thing to do is preprocess the dataset, changing the category value of the symptom attribute and the class attribute with the number 1 and the number -1. The process of extracting raw data obtained in the previous process stage is data that is further processed by data mining methods or used as learning machine learning data at the modeling stage. So that machine learning can predict. The next stage is the evaluation stage, namely, knowing the predictive reliability of the data mining or machine learning method. Then the last stage is the deployment stage to disseminate research results so that they are helpful for implementation by various parties, especially hospitals and other professionals, in the form of developing application programs and scientific articles.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected from medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. There are thousands of data on Covid-19 patients. The patient dataset containing incomplete data and non-covid-19 was omitted or ignored. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

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4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 2 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE II. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses. There were 117 patients with Covid-19 who had a different diagnosis from the others. In this study, the number of signs, symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 3).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or Gi where i = 1, 2, 3 ...12) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 4. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or symptoms.

Furthermore, the preprocessing of the data set is done by changing the Gi with xi and the Gi Yes attribute value with the number 1 while the Gi No attribute with the number -1. In addition, dataset preprocessing is also carried out on class attributes, namely changing the independent isolation class attribute category with the number 1 and the inpatient class attribute category with the number -1, as shown in Table 5.

TABLE III. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE IV. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	Yes	No	No	No	No	No	Inpatient
2	Yes	Yes	No	Yes	No	No	Inpatient
3	Yes	No	Yes	No	No	No	Inpatient
4	Yes	No	No	No	No	No	Inpatient
5	Yes	No	No	No	No	No	Inpatient
6	Yes	No	No	No	Yes	Yes	Inpatient
7	No	No	Yes	No	No	No	Inpatient
...
...
114	No	No	No	No	No	Yes	Self-isolation
115	Yes	No	No	No	Yes	No	Inpatient
116	Yes	No	No	No	Yes	Yes	Inpatient
117	No	No	No	No	No	No	Inpatient

TABLE V. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
...
...
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

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The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, known various programming language [44], which has their respective advantages in building application programs [45][46]. The application program built in this research uses the Python programming language to facilitate patient care status prediction (as show in Table 6).

TABLE VI. USE OF HYPERPARAMETER ON SVM AND RF METHOD

Classifier	Hyperparameter	Value	
SVM	Method		
	Kernel	Rbf	
	Degree	3	
	Gamma	Scale	
	Coef	0	
	Tol	0,000	
	Max_iter	1	
	Random Forest	n_estimators	100
		Criterion	Gini
		Max_depth	None
	Min_samples_split	2	
	Min_samples_leaf	1	

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

In SVM, there are two implementation models: mathematical programming techniques and kernel functions. This study applies kernel functions and focuses on classifying two categories of class attributes. The class attribute is a treatment for $y_i = +1, -1$. The formula of the SVM data mining method is: (a) to form a linear equation from the training data; (b) find the value of w and b, and (c) the value of the classification decision is as follows.

$$((x_1, y_1), \dots, (x_l, y_l)) \quad (1)$$

$$y_i ((w \cdot x_i) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

RF uses bootstrap samples from training data to create a tree from a randomly selected subset. The chosen predictor is a candidate for splitting the decision tree.

The results of the category predictions from the treatment class based on the results of the highest voting were chosen as the final prediction results. The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_{i=1}^c p_i^2 \quad (4)$$

$$Entropy(S) = \sum_{i=1}^c -p_i \times \log_2(p_i) \quad (5)$$

Description:

S = Set of cases

p_i = the proportion of case i to the Set of cases

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and Recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

Accuracy states the closeness of the measurement results to the actual value, while Precision shows how close the difference in the measurement results is on repeated measurements. On the other hand, Recall states the level of success in retrieving information. Precision and Recall are necessary because Precision denotes a measure of quality, and Recall denotes a measure of quantity.

Measurement of accuracy is based on the ratio between the correct predictions (positive and negative) with the overall data. In contrast, precision measurements are based on the percentage of true positive predictions compared to overall positive predicted outcomes. Meanwhile, the recall measurement is based on the ratio of true positive predictions compared to the general actual positive data.

The format of the confusion matrix table is as shown in Table 7. The results of the predictions of the SVM

and RF methods are shown in Tables 8 and 9.

TABLE VII. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

TABLE VIII. CONFUSION MATRIX OF SVM

		Prediction	
		Class	Self-isolation Inpatient
Actual	Self-isolation	4	0
	Inpatient	2	12

TABLE IX. CONFUSION MATRIX OF RF

		Prediction	
		Class	Self-isolation Inpatient
Actual	Self-isolation	4	0
	Inpatient	1	13

4.4.4 K-fold Cross-Validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation helps assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold value. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each subgroup to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the training set (Dtrain). In the second fold, the second subset is the validate set, the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 1 and Figure 3. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE X. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TESTING DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

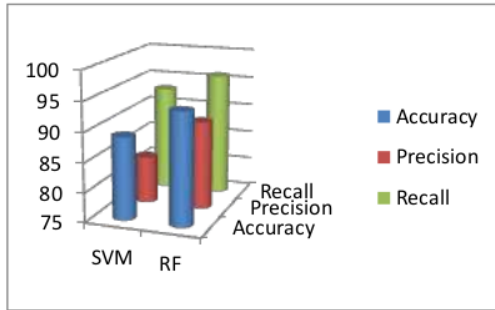


Figure 3. System Model Performance Testing with 85% of Training Data and 15% of Testing Data

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% testing data and 90% training data and 10% testing data are as shown in Table 11.

TABLE XI. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)		Accuracy (in %)		Precision (in %)		Recall (in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 12 and Figure 4.

TABLE XII. PREDICTION PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

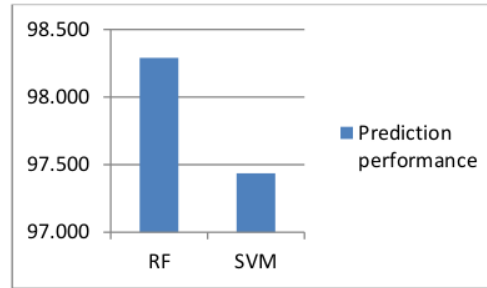


Figure 4. Predictive performance testing using K-fold cross-validation

4.6 Deployment

One of the deployments in this research is making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Thus, the results obtained can be developed and become the knowledge of many parties as a responsibility for the correctness of the effects of research carried out as professional researchers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 90%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%. The research result implies that RF machine learning can help or replace the role of medical personnel in predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, with high accuracy.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Further research needs to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion for future study is: to conduct further research using other data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients and various other diseases, to build a system that not only predicts but also performs clustering, association, and estimates of various other fields of science, including patients' care status, with a combination of

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first, second, and so on as the fifth author.

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Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

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Abstract—Covid-19 has become a global pandemic that causes many deaths, so medical treatment for Covid-19 patients gets special attention, whether hospitalized or self-isolated. However, the problem in medical action is not easy, and the most frequent mistakes are due to inaccuracies in medical decision-making. Meanwhile, machine learning can predict with high accuracy. For that, or that's why this study aims to propose a data mining classification method as a machine learning model to predict the treatment status of Covid-19 patients accurately, whether hospitalized or self-isolated. The data mining method used in this research is the Random Forest (RF) and Support Vector Machine (SVM) algorithm with Confusion Matrix and k-fold Cross Validation testing. The finding indicated that the machine learning model has an accuracy of up to 94% with the RF algorithm and up to 92% with the SVM algorithm in predicting the Covid-19 patient's treatment status. It means that the machine learning model using the RF algorithm has more accurate accuracy than the SVM algorithm in predicting or recommending the treatment status of Covid-19 patients. The implication is that RF machine learning can help/replace the role of medical experts in predicting the patient's care status.

Index Terms—Data Mining, Random Forest, Support Vector Machine, Prediction, Covid-19, Machine Learning

I. INTRODUCTION

The Covid-19 disease is currently a world pandemic [1][2][3]. It is the cause of the global health crisis [4][5][6], which is not only because of its high-speed transmission [5][7], but more than 100 million people have died infected worldwide, and more than two million people have died from it [5]. Covid-19 is a highly contagious viral disease that requires special care and follow-up predictive analytics for better treatment of the disease [8]. However, the Covid-19 pandemic poses a significant challenge to providing health care and services for patients [7]. So it is not surprising that researchers use many research methods to control the Covid-19 pandemic, including the research methods that have received the most attention: prediction, statistical, and epidemiological [6]. Generally, medical actions taken for Covid-19 patients are isolated [9], namely hospitalization or self-isolation. However, these

hospitalized Covid-19 patients are receiving intensive medical care from doctors.

In essence, the care status for Covid-19 patients is self-isolation for patients with non-severe illness status and hospitalization for patients who are seriously ill and at critical risk or cause death. Hospitals or medical doctors take various ways to reduce the number of deaths of Covid-19 patients, including by regulating the status of services in hospitals. Expert doctors recommend that patients self-isolate or should be hospitalized after analyzing the patient's medical data. Determining the level of care for Covid-19 patients is a form of medical treatment or treatment for Covid-19 patients to get proper treatment or care.

Errors in decision-making often occur because decision-makers consider several criteria as the basis for decision-making [10]. So it is not surprising that previous researchers emphasized that the errors most often occur due to inaccuracies in decision making [11], and decision making is a difficult task because of the impact of the decisions made [11]. Likewise, in recommending whether a Covid-19 patient should be hospitalized or self-isolated, several criteria from the disease symptoms and the results of medical tests are the basis for considering whether a patient should be hospitalized or self-isolated. In essence, it is difficult to accurately determine the treatment status of Covid-19 patients, both inpatient and self-isolation.

Meanwhile, Machine Learning is a rapidly growing part of computer science today [12]. Although in most scientific studies, machine learning is popular, it is still very limited in health studies [13]. Machine learning helps mining data to predict mining results accurately [14]. Machine learning is a helpful technique for finding correlations based on cases to predict [15]. With the availability of big data, it is possible to develop various solutions using machine learning [16][17]; moreover, with advances in information and communication technology [18][19], it is straightforward to collect the required big data. Among the solutions using machine learning, one of which is predictive modeling [20][21][22]. Furthermore, machine learning can uncover hidden patterns in big data, distinguish patterns better and more accurately [13], and provide high-accuracy prediction results [23]. For this reason (or

why), this study's objective is to propose a machine learning system model for decision-making solutions (predictions) for the treatment status of Covid-19 patients, both inpatient and self-isolation, using data mining methods.

The implication is that the proposed machine learning system model can help and even replace the role of medical experts (specialist doctors) in making medical decisions for Covid-19 patients, whether hospitalization or self-isolation. Machine learning performs tasks like a medical specialist in deciding the results of the diagnosis of the nursing status of Covid-19 patients based on medical data of Covid-19 patients. Furthermore, machine learning can work tirelessly, time and place, and has intelligence like an expert, so it is not surprising that previous research confirms that intelligent machines can make superior decisions to experts because humans have a human error factor [24].

Machine learning can predict classification to predict class membership and regression to show numerical values [12]. While data mining is part of machine learning that can make system models have artificial intelligence. Artificial intelligence is a breakthrough in today's technology that has been widely used in prediction [25]. The embodiment of artificial intelligence in machine learning with data mining methods is an iterative process of training and repeated testing of data sets (big data) on the system model. In short, machine learning has an artificial intelligence role in predicting new data with high accuracy [23]. After all, predicting individuals with symptoms of being infected with Covid-19 mandates machine learning (application-based) and contributes to effectively isolating Covid-19 patients [26].

Big data demands large storage media [27]. However, big data is no longer traditionally processed [28]. Instead, today's big data processing relies on machines that can provide systematic results [29]. Big data storage is generally on a computer server with a large storage capacity. Still, some also make it happen by renting online cloud data storage services such as Amazon Simple Storage Service (Amazon S3) and Google Cloud [30]. Cloud facilitates cost-effective big data storage and analysis [30].

Big data processing techniques in data mining include several stages: target data, preprocessed data, data mining, and evaluation/analysis of mining results [30]. Target data and preprocessed data are the processes of extracting raw data from big data [30]. Target data is to select the required data (sample data) and classify data. The preprocessed data is to prepare data sets for data mining, including cleaning up incomplete data, duplicate data, and converting string data into numeric coded data. Finally, data mining and evaluation results extract hidden information by applying data mining methods suitable for the objectives and analyzing them.

Many Data Mining methods include K-Means, Naïve Bayes, KNN (k-Nearest Neighbor), ID3 (Iterative

Dichotomiser 3), C4.5, Cart, RF, SVM, and others. There are two types or methods of machine learning, namely supervised machine learning and unsupervised machine learning. It is referred to as a supervised learning method when the subject's membership is known, and training is carried out to classify new data into its category. On the other hand, it is referred to as an unsupervised learning method when the subject's membership is unknown, and the closest distance search is to categorize the groups. The Data Mining methods used in this research are RF and SVM algorithms. RF and SVM are prevalent machine learning algorithms used in various scientific studies [31] and constitute data classification techniques with supervised learning methods. The RF machine learning algorithm has been widely applied for classification [32] [33], as well as SVM algorithm has a widely known technique used for classification [33]. It is why this study uses SVM and RF to classify treatment status. Given that the SVM and RF machine learning algorithms are both popularly used by many researchers, the SVM and RF machine learning algorithms are the most appropriate combination used in research, including research to classify and predict the treatment status of Covid-19 patients.

However, it is essential to know the accuracy of predicting the care status of Covid-19 patients from the system model proposed in this study and whether the patient should be hospitalized or self-isolated. Therefore, this study also further tested the percentage of machine learning efficacy or accuracy in predicting the treatment status of Covid-19 patients. The accuracy of predicting the treatment status of Covid-19 patients is tested on both RF and SVM machine learning methods.

The organization of the following writing of this manuscript is as follows: The second subsection discusses several of the related works of previous researchers and their relevance to the work in this research article. The third subsection describes Research Methodology, which discusses methods used in research in recommending patient care status. Meanwhile, the fourth subsection discusses the results of the study. Finally, it ends with a subsection that concludes the study's findings, the novelty of the research results, and advice for further research.

II. RELATED WORKS

This subsection provides an overview of some related works from the latest scientific articles compared with the work in this research article.

Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) performed the classification of satellite remote sensing images using machine learning algorithms with RF, SVM, and decision tree classifier (DT) [31]. This previous research is different from the research in the article on the research objectives and the object under study. In the meantime, Celestine Iwendi et al. (2020) proposed the Random Forest model to predict the disease severity of Covid-19 patients [34]. The

difference between previous research and the research in this article is that the previous research only used one method, namely Random Forest. In contrast, the research in this article used two methods, namely Random Forest and SVM. The difference also lies in the prediction criteria and class; previous research predicts the severity of the illness of Covid-19 patients, while the research in this article predicts the treatment status of Covid-19 patients.

Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) proposed a machine learning model to classify breast cancer by applying RF and SVM methods [35]. Previous research and the article in this research are both using RF and SVM methods. However, the previous research has research objectives that are not the same as the research in this article. The prior study classified breast cancer as patients with breast cancer. In contrast, the research in this article predicts the treatment status of Covid-19 patients.

Based on patient clinical data, using statistical methods, Boran Hao et al. (2020) developed a model to predict pneumonia severity in Covid-19 patients using the Natural Language Processing tool [36]. However, this previous research differs in the research purpose and way compared to the research in this article. Meanwhile, Anthony Anggrawan et al. (2021) implemented machine learning to diagnose drug users and types of drug-using Forward Chaining and Certainty Factor methods [23]. Meanwhile, the research in this article develops machine learning to predict the patient's treatment status, whether inpatient or self-isolation, based on symptoms or patient medical data using RF and SVM.

Hongwei Zhao et al. (2021) built a model to predict the number of cases of Covid-19 patients in the future using the Poisson and Gamma distribution [37]. Similarities between articles in this study and the previous one proposed a model with a machine learning approach. However, this previous research differs in the research purpose and method compared to the research in this article. In the meantime, Bassam Mahboub et al. (2021) developed a model to predict the length of hospital stay with the decision tree (DT) method [8]. This article's research differs from previous research; the difference lies in the research objectives and techniques used. If prior research predicts the length of stay for Covid-19 patients, the research in this article predicts whether Covid-19 patients should be hospitalized or self-isolated. This article's research does not use the DT method but uses the RF and SVM methods.

Soham Guhathakurata et al. (2021) predicted whether a person is infected with Covid-19 or not using SVM [38]. However, this previous study differed in its objectives from this article's research. The previous research predicts patients suffering from Covid-19 or not utilizing the SVM data mining method. In contrast, the research in this article indicates the patient's care status using RF and SVM data mining methods. At the

same time, Ankit Mehrotraa and Reeti Agarwal (2021) reviewed the usefulness of the Data Mining method for the Covid-19 pandemic [39]. This previous research is a literature review study that concludes that the Data Mining method plays an essential role in health care, diagnosing diseases, and recommending cures. However, it is different from this article's research because it is an experimental study, not a literature review.

Pratiyush Guleria et al. (2022) proposed a machine learning model to predict the death rate of Covid-19 patients [14]. However, previous research has different objectives and data mining methods compared to this article's research. The difference is that previous studies examined the infection rate of Covid-19 patients to predict the cure/death rate of Covid-19 patients using the SVM, Decision Trees, and Naïve Bayes data mining methods. In contrast, the research in this article predicts the care status of Covid-19 patients using RF and SVM data mining methods [14].

Anthony Anggrawan, Mayadi, Christofer Satria, and Lalu Ganda Rady Putra (2022) developed a machine learning model for scholarship recipients' recommendations by using Analytical Hierarchy Process (AHP) and the Multi-Objective Optimization Method by Ratio Analysis (Moora) methods [40]. However, the previous research differs in the purpose and way compared to this article's research. In contrast, Vadim Demichev et al. (2022) offered a model to optimize the treatment or intensive care of seriously ill Covid-19 patients with plasma proteomics [41]. This previous research is different from the research in the article on the research objectives, research method, and the object under study.

Table 1 compares some of the most recent previous related work with the work carried out in this study. By referring to the elaboration of the most recent last related work by some researchers, the research carried out in this article has novelties (from the prior research gap) that previous researchers have not studied. In essence, the gap in earlier research is that no one has researched machine learning models to predict the inpatient status or self-isolation of Covid-19 patients by involving RF and SVM algorithms. In addition, the 12 criteria used to indicate the treatment status of Covid-19 patients are entirely different from previous similar studies (as shown in Table 1 in the Criteria/Attributes column). So, the study's originality lies in proposing a machine learning model to predict the nursing status of Covid-19 patients, whether inpatient or self-isolation, which previous researchers have never done. Besides that, the novelty is also in the method used, not just one data mining method in predicting the treatment status of Covid-19 patients, but using two data mining methods. So this study can show differences in the accuracy of the RF and SVM methods in predicting the treatment status of Covid-19 patients.

TABLE I. COMPARISON OF THIS ARTICLE'S WORK WITH SOME PREVIOUS RELATED WORKS

Research by	Research methods			Criteria/Attributes		Research Object	Accuracy Test
	RF	SVM	ML	Number	Name		
Askin Kavzoglu, Furkan Bilucan, and Alihan Teke (2020) [31]	Yes	Yes	Yes	10	Coastal Aerosol, Blue, Green, Red, Vegetation Red Edge, NIR, Narrow NIR, Water vapor, SWIR-Cirrus, SWIR	Satellite remote sensing images	Yes
Celestine Iwendi et al. (2020) [34]	Yes	No	Yes	6	Symptom1, symptom2, symptom3, symptom4, symptom5, symptom6	Illness severity of Covid-19 patients	No
Chelvian Aroef, Yuda Rivan, and Zuherman Rustam (2020) [35]	Yes	Yes	Yes	9	Age, Body Mass Index (BMI), Glucose, Insulin, Homa, Leptin, Adiponectin, Resistin, MCP 1	Breast cancer	Yes
Boran Hao et al. (2020) [36]	No	No	No	10	Radiology Opacities, Respiratory Rate, Age, Fever Male, Albumin, Anion Gap, SpO2, LDH, Calcium	The severity of pneumonia in Covid-19 patients	Yes
Anthony Anggrawan et al. (2021) [23]	No	No	Yes	27	Out of breath, Anxious, Nausea, Diarrhea, Convulsions, Easily angry, Depression, Sleep patterns change, Sweating, Chills, Shaking, Insomnia, Fast heart rate, Blood pressure rises, Difficult to focus, Difficult to rest, Weight loss, Dry mouth, Blurred vision, Changed skin color, Constipation, Stomachache, Drowsiness, Itching, Difficulty urinating, Mood swings, Dizziness	Drug users and types of drug-using	Yes
Hongwei Zhao et al. (2021) [37]	No	No	No	0	-	Number of cases of Covid-19 patients	No
Bassam Mahboub et al. (2021) [8]	No	No	Yes	5	Age, Gender, Nationality, Blood group, BMI	The treatment period for Covid-19 patient	Yes
Soham Guhathakurata et al. (2021) [38]	No	Yes	Yes	8	Temp, Breathing rate, Hypertension, Heartbeat rate (HBR), Acute respiratory disease syndrome (ARDS), Chest pain, Heart disease, Cough with sputum (CWS)	Predicting whether patients are infected with Covid-19 or not	No
Ankit Mehrotraa and Reeti Agarwal (2021) [39]	No	No	No	0	-	Discussing the data mining method's role in the Covid-19 pandemic	No
Pratiyush Guleria et al. (2022) [14]	No	Yes	Yes	0	-	The death rate of Covid-19 patient	Yes
Anthony Anggrawan, et al. (2022) [40]	No	No	Yes	6	Achievement index, achievement points, recommendation, organizational activity, semester level, and completeness of documents	Scholarship recipient	Yes
Vadim Demichev et al. (2022) [41]	No	No	No	0	-	Optimization of treatment for Covid-19 patients	Yes
Our research	Yes	Yes	Yes	12	Pneumonia, ARDS, CHF, AKI, CAD, Dyspnea, NSTEMI, ADHF, HHD, Febris, Anosmia, Ageusia	Care status of Covid-19 patients, whether inpatient or self-isolation	Yes

Note: ML = Machine Learning

III. METHODOLOGY

This study applies two data mining methods or machine learning algorithms: RF and SVM. The big data is on Covid-19 patients from a regional hospital in Mataram, Indonesia. The significant data source used in this study is primary data from patient medical records/documents. The attributes of the patient's disease symptoms and care status classes amount to thousands of patient medical record data. Patient datasets containing non-Covid-19 and duplicate and incomplete Covid-19 patient data are removed, so only data is left as a dataset for data mining processes. Medical record data of disease symptoms obtained

from string data is then converted into numeric data. The development of the application program in this study uses the Python computer programming language.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods. The data mining process in this research uses CRISP-DM (Cross-Industry Standard Process for Data Mining). CRISP-DM is a standard data mining process. The process in CRISP-DM comprises a six-stage [42], as shown in Figure 1 [43]. Figure 2 shows the process carried out at each stage of CRISP-DM.

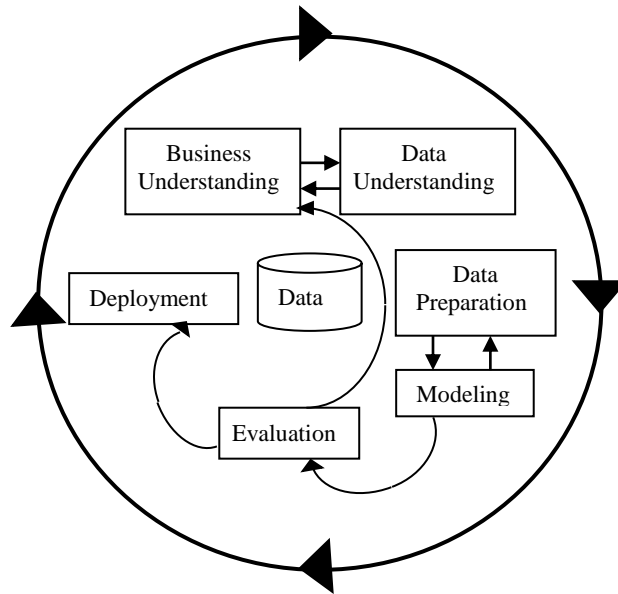


Figure 1. The CRISP-DM Process

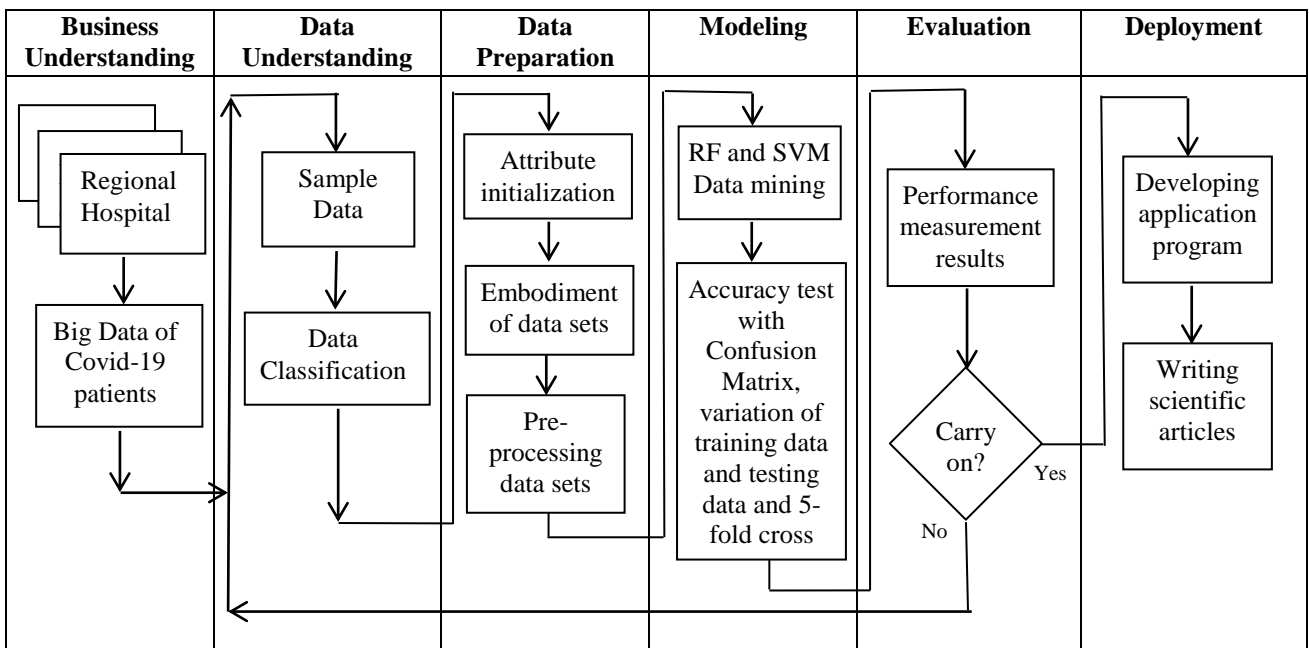


Figure 2. Data Mining Process of Covid-19 Patient Big Data

In Figure 2, business understanding is the stage of sorting out thousands of hospital patient medical data to collect the required patient data. The next stage is understanding the data collected as representative data for Covid-19 patients. This Covid-19 patient data classifies the patient's signs and symptoms and treatment status, which needs further processing at the next stage. The next stage is the data preparation stage, which essentially determines the attributes of the names of signs and symptoms of Covid-19 patients. The embodiment of the dataset containing knowledge according to treatment status refers to the signs and symptoms that the patient has (marked with Yes) or does not have (marked with No). The next thing to do is preprocess the dataset, changing the category value of the symptom attribute and the class attribute with the number 1 and the number -1. The process of extracting raw data obtained in the previous process stage is data that is further processed by data mining methods or used as learning machine learning data at the modeling stage. So that machine learning can predict. The next stage is the evaluation stage, namely, knowing the predictive reliability of the data mining or machine learning method. Then the last stage is the deployment stage to disseminate research results so that they are helpful for implementation by various parties, especially hospitals and other professionals, in the form of developing application programs and scientific articles.

This research uses a confusion matrix and k-fold cross-validation to measure the classification performance of RF and SVM methods.

IV. RESULT AND DISCUSSION

4.1 Business Understanding

The significant data acquisition of Covid-19 patients needed for research is obtained from the hospital. The data collected from medical document data from all Covid-19 patients registered at the hospital includes the patient's name, disease symptoms, and the treatment status specified. There are thousands of data on Covid-19 patients. The patient dataset containing incomplete data and non-covid-19 was omitted or ignored. The critical information extracted at this stage is first to find the attributes or criteria of the class of treatment status (hospitalization or self-isolation); second, to find the category of each feature of the treatment status class. The existing attributes and categories represent disease symptoms and other medical data from Covid-19 patients. Based on Covid-19 patient data adopted from the hospital, there are 12 symptom criteria or patient medical data that are used as references by expert doctors in determining the status of patient care, whether to be hospitalized or self-isolated. Furthermore, big data containing several symptom criteria or patient medical data is used in training and testing the prediction model proposed in this study. Therefore the offered machine learning model has artificial intelligence in predicting.

4.2 Data Understanding

The Data Understanding stage is preparing the data set from the research. The dataset from this study is a data representation of the Covid-19 patient sample, which contains sign and symptom data and treatment status. Table 2 shows the association between signs and symptoms of disease and treatment status in the study data set.

TABLE II. DATA SET OF THE SIGNS AND SYMPTOMS AND THE TREATMENT STATUS OF COVID-19 PATIENTS

No	Disease Sign and Symptom	Treatment
1	Pneumonia, Dyspnea	Inpatient
2	Pneumonia, ARDS, AKI, Febris	Inpatient
3	Pneumonia, CHF, CAD, Dyspnea	Inpatient
4	Pneumonia, AKI	Inpatient
5	Pneumonia, CAD	Inpatient
6	Pneumonia, Dyspnea, Anosmia, Ageusia	Inpatient
7	CHF, NSTEMI	Inpatient
..
..
114	Febris, Anosmia, Ageusia	Self-isolation
115	Pneumonia, Anosmia	Inpatient
116	Pneumonia, Anosmia, Ageusia	Inpatient
117	HHD	Inpatient

4.3 Data Preparation

Each patient confirmed positive for Covid-19 has a different diagnosis from others, and some patients have similar diagnoses. There were 117 patients with Covid-19 who had a different diagnosis from the others. In this study, the number of signs and symptoms or the number of research criteria is 12 signs and symptoms, or the number of research criteria is 12 signs and symptoms or 12 criteria (see Table 3).

The signs and symptoms of each Covid-19 patient (G01, G02, ... G12 or G_i where $i = 1, 2, 3 \dots 12$) are not all the same from one patient to another. For this reason, the attributes of each patient's data are different, and some are the same between one patient and another, as shown in Table 4. If the sign or symptom attribute is No, the patient does not have these signs or symptoms. On the other hand, if the sign or symptom attribute is Yes, the patient has these signs or symptoms.

Furthermore, the preprocessing of the data set is done by changing the G_i with x_i and the G_i Yes attribute value with the number 1 while the G_i No attribute with the number -1. In addition, dataset preprocessing is also carried out on class attributes, namely changing the independent isolation class attribute category with the number 1 and the inpatient class attribute category with the number -1, as shown in Table 5.

TABLE III. DATA SET RELATED TO RESEARCH ATTRIBUTES AND DISEASE SIGNS AND SYMPTOMS

Attribute	Sign and Symptoms	Word extension
G01	Pneumonia	Pneumonia
G02	ARDS	Acute Respiratory Distress Syndrome
G03	CHF	Congestive Heart Failure
G04	AKI	Acute Kidney Injury
G05	CAD	Coronary Artery Disease
G06	Dyspnea	Dyspnea
G07	NSTEMI	Non-ST-Segment Elevation Myocardial Infarction
G08	ADHF	Acute Decompensated Heart Failure
G09	HHD	Hypertensive Heart Disease
G10	Febris	Febris
G11	Anosmia	Anosmia
G12	Ageusia	Ageusia

TABLE IV. DATA SET OF KNOWLEDGE BASED ON TREATMENT STATUS REFERRING TO THE SIGNS AND SYMPTOMS

No	G01	G02	G03	G04	G11	G12	Class
1	Yes	No	No	No	No	No	Inpatient
2	Yes	Yes	No	Yes	No	No	Inpatient
3	Yes	No	Yes	No	No	No	Inpatient
4	Yes	No	No	No	No	No	Inpatient
5	Yes	No	No	No	No	No	Inpatient
6	Yes	No	No	No	Yes	Yes	Inpatient
7	No	No	Yes	No	No	No	Inpatient
..
..
114	No	No	No	No	No	Yes	Self-isolation
115	Yes	No	No	No	Yes	No	Inpatient
116	Yes	No	No	No	Yes	Yes	Inpatient
117	No	No	No	No	No	No	Inpatient

TABLE V. PREPROCESSING OF THE DATA SET RESULT

No	G01	G02	G03	G04	G11	G12	Class
1	1	-1	-1	-1	-1	-1	-1
2	1	1	-1	1	-1	-1	-1
3	1	-1	1	-1	-1	-1	-1
4	1	-1	-1	-1	-1	-1	-1
5	1	-1	-1	-1	-1	-1	-1
6	1	-1	-1	-1	1	1	-1
7	-1	-1	1	-1	-1	-1	-1
..
..
114	-1	-1	-1	-1	-1	1	1
115	1	-1	-1	-1	1	-1	-1
116	1	-1	-1	-1	1	1	-1
117	-1	-1	-1	-1	-1	-1	-1

4.4 Modeling

The proposed machine learning model to predict Covid-19 treatment status in this study applies the RF and SVM data mining classification methods. In addition, known various programming language [44], which has their respective advantages in building application programs [45][46]. The application program built in this research uses the Python programming language to facilitate patient care status prediction (as show in Table 6).

TABLE VI. USE OF HYPERPARAMETER ON SVM AND RF METHOD

Classifier	Hyperparameter	Value
SVM	Method	
	C	1
	Kernel	Rbf
	Degree	3
	Gamma	Scale
	Coef	0
	Tol	0,000
		1
	Max_iter	-1
	RF	n_estimators
Criterion		Gini
Max_depth		None
Min_samples_split		2
Min_samples_leaf		1

4.4.1 SVM data mining method

The process of realizing the classification using the SVM data mining method is as follows: (a) Forming a linear equation from the training data that has gone through the preprocessing stage; (b) Finding the values of w and b by means of elimination and substitution of linear equations; (c) Finding the value of the classification decision with the function.

In SVM, there are two implementation models: mathematical programming techniques and kernel functions. This study applies kernel functions and focuses on classifying two categories of class attributes. The class attribute is a treatment for $y_i = +1, -1$. The formula of the SVM data mining method is: (a) to form a linear equation from the training data; (b) find the value of w and b , and (c) the value of the classification decision is as follows.

$$S = ((x_1, y_1), \dots, (x_l, y_l)) \quad (1)$$

$$y_i ((w \cdot x_i) + b) \geq 1, i = 1, \dots, \quad (2)$$

$$(x) = w \cdot x + b \quad (3)$$

Description:

S= set; x = attribute; y = class; w = weight; b = bias

4.4.2 RF data mining method

The process of realizing the classification using the RF data mining method is as follows: (a) Generating a random subset of data; (b) Creating a decision tree (Root tree, branch tree & leaves tree) from each attribute and class; and (c) Testing each decision tree with data testing and calculating the accuracy of each decision tree.

RF uses bootstrap samples from training data to create a tree from a randomly selected subset. The chosen predictor is a candidate for splitting the decision tree.

The results of the category predictions from the treatment class based on the results of the highest voting were chosen as the final prediction results. The formula for the RF data mining method is the Gini criterion and the Entropy criterion:

$$Gini = 1 - \sum_{i=1}^c p_i^2 \quad (4)$$

$$Entropy(S) = \sum_{i=1}^c -p_i \times \log_2(p_i) \quad (5)$$

Description:

S = Set of cases

p_i = the proportion of case i to the Set of cases

4.4.3 Confusion matrix

This research uses a confusion matrix to measure the performance of the classification method. The confusion matrix is a method that can be used to measure the performance of a classification method. In essence, the confusion matrix can produce information by comparing the system's classification results with the classification results that should be.

In measuring performance using the Confusion Matrix, four terms represent the results of the classification process, namely: True Positive (TP) or positive data detected correctly; False Positive (FP) or negative data detected is positive; True Negative (TN) or negative data detected correctly, and False Negative (FN) or positive data detected is negative. Meanwhile, the calculation of accuracy, prediction, and Recall in the confusion matrix can use the following equation:

$$accuracy = \frac{TP+TN}{TP+TN+FP+FN} \quad (6)$$

$$Precision = \frac{TP}{TP+FP} \quad (7)$$

$$Recall = \frac{TP}{TP+FN} \quad (8)$$

Accuracy states the closeness of the measurement results to the actual value, while Precision shows how close the difference in the measurement results is on repeated measurements. On the other hand, Recall states the level of success in retrieving information. Precision and Recall are necessary because Precision denotes a measure of quality, and Recall denotes a measure of quantity.

Measurement of accuracy is based on the ratio between the correct predictions (positive and negative) with the overall data. In contrast, precision measurements are based on the percentage of true positive predictions compared to overall positive predicted outcomes. Meanwhile, the recall measurement is based on the ratio of true positive predictions compared to the general actual positive data.

The format of the confusion matrix table is as shown in Table 7. The results of the predictions of the SVM

and RF methods are shown in Tables 8 and 9.

TABLE VII. CONFUSION MATRIX

Class	Classified Positive	Classified Negative
Positive	True Positive	False Negative
Negative	False Negative	True Positive

TABLE VIII. CONFUSION MATRIX OF SVM

		Prediction	
		Class	Self-isolation Inpatient
Actual	Self-isolation	4	0
	Inpatient	2	12

TABLE IX. CONFUSION MATRIX OF RF

		Prediction	
		Class	Self-isolation Inpatient
Actual	Self-isolation	4	0
	Inpatient	1	13

4.4.4 K-fold Cross-Validation

This study used K-fold cross-validation to measure the performance of the classification method. K-fold cross-validation helps assess the performance of data mining methods by dividing the data sample randomly and grouping the data as much as the k-fold value. In the performance testing of this study with k-fold cross-validation, the dataset is partitioned into five subsets (k = 5). It allows each subgroup to have the same number and fold, which refers to the number of resulting subsets. Dataset partitioning is done by taking random samples from the dataset. However, data that has been taken previously will not be retrieved.

In the first fold, the first subset serves as the validate set (Dval), and the remaining four subsets serve as the training set (Dtrain). In the second fold, the second subset is the validate set, the remaining subset is the training set, and so on until the 5th fold.

4.5 Evaluation

The evaluation of the proposed model in this study is to measure the performance of the resulting prediction system model. The model's performance evaluation is based on the prediction system model generated by the RF and SVM methods.

4.5.1 Evaluation of prediction model with confusion matrix

Evaluation of the prediction results of the proposed system model uses the confusion matrix technique. The evaluation result using the confusion matrix is shown in Table 10 and Figure 3. The accuracy in predicting with 85% of training data and 15% of test data shows that the RF machine learning method is more accurate and precise than the SVM machine learning method.

TABLE X. SYSTEM MODEL PERFORMANCE TESTING WITH 85% OF TRAINING DATA AND 15% OF TESTING DATA

Method	Accuracy	Precision	Recall
SVM	89%	83%	93%
RF	94%	90%	96%

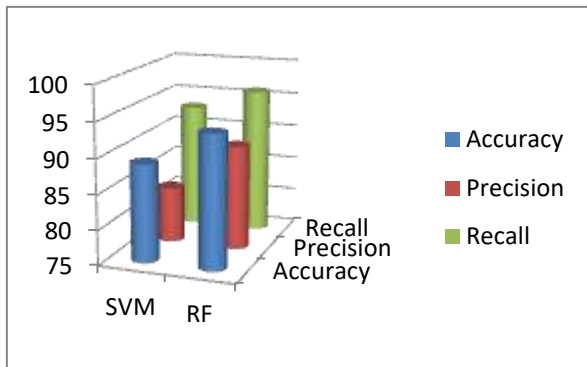


Figure 3. System Model Performance Testing with 85% of Training Data and 15% of Testing Data

Further comparison of the accuracy and precision of the prediction system model with 50% training data and 50% testing data, 60% training data and 40% testing data, 70% training data and 30% testing data, 80% training data and 20% data 90% testing and training data and 10% testing data are as shown in Table 11.

TABLE XI. PREDICTION SYSTEM MODEL PERFORMANCE TESTING WITH VARIOUS TEST DATA AND TRAINING DATA VARIATIONS

Data (in %)		Accuracy (in %)		Precision (in %)		Recall (in %)	
Training	Testing	RF	SVM	RF	SVM	RF	SVM
50	50	95	97	97	96	91	96
60	40	96	91	97	90	93	90
70	30	94	92	96	91	92	90
80	20	96	92	94	89	97	94
90	10	92	83	75	67	95	91
Average		95	91	92	87	94	92

Predicting with various test data and training data variations shows that the RF machine learning method is more accurate and precise than the SVM machine learning method. In other words, the prediction system model proposed to predict the treatment status of Covid-19 patients using the RF method is better (more accurate and precise) than the SVM machine learning method based on performance tests with a confusion matrix.

4.5.2 Evaluation of prediction model with k-fold cross-validation

The performance of the model proposed in this study uses a 5-fold cross-validation on both RF and SVM prediction models presented in Table 12 and Figure 4.

TABLE XII. PREDICTION PERFORMANCE TESTING WITH K-FOLD CROSS-VALIDATION

RF (in %)	SVM (in %)
98.290	97.436

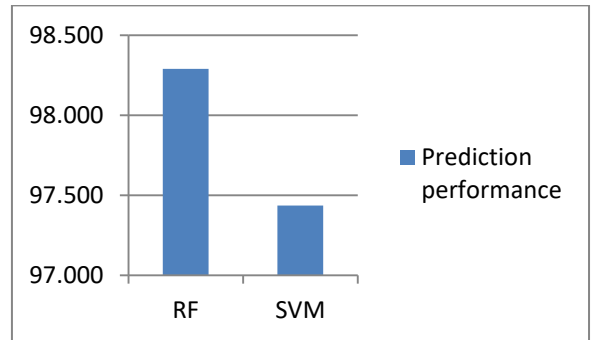


Figure 4. Predictive performance testing using K-fold cross-validation

4.6 Deployment

One of the deployments in this research is making scientific articles on machine learning system models that are produced to be published in reputable scientific papers. Thus, the results obtained can be developed and become the knowledge of many parties as a responsibility for the correctness of the effects of research carried out as professional researchers. Another form of deployment is to make reports to cooperative hospital partners where data on Covid-19 patients is obtained.

V. CONCLUSION

This study found that the prediction system model for the treatment status of Covid-19 patients using the RF machine learning method had better predictive performance than the SVM machine learning method. The test of accuracy and precision in predicting the treatment status of Covid-19 patients using the confusion matrix showed that the RF machine learning method has a prediction accuracy of 94% and a precision of 90%; In comparison, the SVM machine learning method has a prediction accuracy of 89% and a precision of 83%. Further testing of the accuracy of the system model in predicting the treatment status of Covid-19 patients using k-fold cross-validation showed that the RF machine learning method had a prediction accuracy of 98.290% and the SVM machine learning method had a prediction accuracy of 97.436%. The research result implication is that RF machine learning can help or replace the role of medical personnel in predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, with high accuracy.

The novelty of this study is to propose a system model for predicting the treatment status of Covid-19 patients, whether inpatient or self-isolation, which researchers have never studied before using two machine learning methods of RF and SVM.

Further research needs to develop a machine learning system model to predict the death or recovery status of each Covid-19 patient. Another suggestion for future study is: to conduct further research using other data mining methods to predict patient care status and the status of death or recovery from Covid-19 patients and various other diseases, to build a system that not only predicts but also performs clustering, association, and estimates of various other fields of science, including patients' care status, with a combination of

machine learning and the Internet of Things.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors undertake work assignments to complete the research and writing of this article jointly. The level of roles and tasks of research work is the basis that places each author as the first, second, and so on as the fifth author.

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Dear Anthony Anggrawan, Mayadi, Christofer Satria, Krismono, Ria Rismayati:

We are pleased to inform you that the following paper has been officially accepted for publication in Journal of Advances in Information Technology.

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Title: Comparative Analysis of Machine Learning in Predicting the Treatment Status of Covid-19 Patients

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