

## **The impact of imputation procedures with machine learning methods on the performance of classifiers: An application to coronary artery disease data including missing values.**

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### **Abstract**

**Prediction and learning in the presence of missing data are pervasive problems in data analysis by machine learning. This study focuses on the problems of collaborative classification with missing data on Coronary Artery Disease (CAD) and suggests alternative imputation methods in the case of the lack of laboratory test as well other specific parameters. This study develops three novel data imputation methods utilizing machine learning algorithms (K-means, Multilayer Perceptron (MLP), and Self-Organizing Maps (SOMs)) and compares the performance of our methods with well-known mean method. Benchmark classification methods (Logistic Model Trees (LMT), MLP, Random Forest (RF), and Support Vector Machine (SVM)) are used to conduct experiments on CAD dataset after imputation. The performance of the classifiers is evaluated according to the values of accuracy, specificity, sensitivity, f-measure, precision and normalized root mean square error. Based on statistical analysis, the SOM imputation method achieves the best values for accuracy (88.23%), F-measure (0.879), and precision (0.881). Moreover, MLP is mostly more stable than other imputation methods when the mean scores of the results of classifiers are considered. According to the results, the data imputation experiments conducted in this study suggests that machine learning imputation methods increase the prediction performance of the classifiers and strengthen disease-diagnosed success.**

**Keywords:** Missing value, Self-organizing maps, Multilayer perceptron, K-means, Coronary artery disease, Classification methods.

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### **Introduction**

Acute chest pain is one of the frequent causes of presentation to emergency department; however, only 15-20% these presentations of chest pain really have [1]. In addition, it becomes difficult to diagnose the ones which do not have specific symptoms or electrocardiographic signs. Furthermore, there are many factors which cause higher risk for CAD and can be classified according to the National Cholesterol Education Program (NCEP) ATP III Guidelines.

Hospital Information Systems (HISs) are used to collect and provide health information for decision-making and research in hospitals. Studies conducted in the medical field suggest that datasets are formed manually in some cases while they are formed through retrieval from HISs in others [2]. In the records of datasets formed by manual data entry in clinical studies, there is a striking amount of missing values (MVs). Therefore, it is very important to impute MVs for application to medical

datasets, and this is an indispensable component of the pre-processing stage in classification problems [3].

The imputation method is one of the tools used to solve MV problems in medical data analysis, and its application depends on the distribution of MVs in the data. The mean/mode imputation method is an example of transforming MVs into new continuous and categorical values depending on the distribution of their features. Recently, fuzzy unordered rule induction algorithm imputation and machine learning have been used as alternative MV imputation methods for many clinical datasets and others with MVs [4-6]. In the literature, machine learning methods have better performances when compared to traditional methods. The methods used in the studies of MV imputation with machine learning in many different domains are Multilayer Perceptron (MLP) [7], Self-Organizing Maps (SOMs) [8,9], Decision Tree (DT) [10] and K-Nearest Neighbors (KNN) [5]. Furthermore, not all algorithms are suitable for all MV problems; hence, it is

necessary to choose a suitable method depending on the nature of the study and the structure of the dataset.

This study aims to make more stable diagnosis predictions on the CAD dataset after imputation procedures, and the imputation of MV problem has been examined with the machine learning approach as an alternative to the traditional mean method.

## Material and Methods

### Description of cardiovascular dataset

This study included 459 patients who presented to the department of cardiology of Mersin Research and Training Hospital, Mersin, Turkey with the suspect of CAD. Angiography is applied as a standard procedure to determine the stenosis. After angiography has been used to determine the location of the lesion, disease of the left main coronary artery (stenosis diameter > 50% of vessel diameter) is considered to indicate the presence of CAD. The present study categorizes 22 clinical comorbidities, cardiac status, medical history features into categorical-type and continuous-type format shown in Table 1.

**Table 1.** Clinical characteristics relevant to diagnosis missingness percentages are shown for numeric features which have MVs.

Feature label	Variable type	Missingness (%)
Age (Individual's age)	Quantitative/numeric	-
Gender (Individual's gender)	Qualitative/categorical	-
DM (Diabetes mellitus)	Qualitative/categorical	-
FH (Family history)	Qualitative/categorical	-
HTN (Hypertension)	Qualitative/categorical	-
Smoke	Qualitative/categorical	-
Hyp (Hyperlipidemia)	Qualitative/categorical	-
LDL (Low-Density Lip.) (mg/dl)	Quantitative/numeric	13.07
HLD (High-Density Lip.) (mg/dl)	Quantitative/numeric	12.41
TG (Triglyceride) (mg/dl)	Quantitative/numeric	12.41
TC (Total Cholesterol) (mg/dl)	Quantitative/numeric	13.72
Urea (mg/dl)	Quantitative/numeric	16.78
Hb (Hemoglobin) (g/dl)	Quantitative/numeric	1.08
Cr (Creatine) (mg/dl)	Quantitative/numeric	0.6
RDW (Red cell distribution) (mg)	Quantitative/numeric	3.7
MCV (Mean corpus. volume) (mg)	Quantitative/numeric	3.7
EF (Electrocardiographic result) (%)	Quantitative/numeric	18
FBS (Fasting blood sugar) (mg/dl)	Quantitative/numeric	5.22
Asthma	Qualitative/categorical	-
Renal failure	Qualitative/categorical	-

COPD	Qualitative/categorical	-
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The cardiovascular dataset has 314 cases of CAD and 145 cases without CAD with 22 features for each case. Of the total number of cases in the dataset, 277 are completed with all the features and 182 have at least one MV. The complete part of the dataset consists of 210 CAD and 67 without CAD samples. We observed that 182 out of 459 cases have 4.8% to 33.3% MVs in their features. The CAD dataset contains 11 continuous-type of features which has MV at random drop-outs.

### Imputation techniques

The dataset is separated into two subsets, where the first subset (referred to as set 1) consists of cases that do not include MVs and the second subset (referred to as set 2) has cases that include MVs. The cases in set 2 are listed in ascending order depending on their numbers of MV features. The k-fold cross validation was used in this work to ensure that all data points are used for both training and validation. There are some of the notations as shown in Table 2 placed in imputation algorithms.

**Table 2.** Table of notations.

Symbol	Description
$N_x$	MVs sample
M1	Complete part of the CAD dataset (set 1)
M2	The part of the CAD dataset has MVs (set 2)
$C_i$	The $i^{\text{th}}$ cluster
$c_i$	The centroid of cluster $C_i$
T	Target vector
W	Wight vector of BMN
$\Theta$	Restraint due to the distance of BMN

### K-means imputation process

The K-means cluster method partitions the n samples into k  $\leq$  n sets  $S=(S_1, S_2, \dots, S_k)$  so as to minimize the variance of the within-cluster sum of squares [11]. The number of clusters is assigned considering the similarity between the cluster centres. For set 1, the number of clusters k is chosen as 6. The algorithm used to carry out the MV imputation method by using the K-means is as follows.

#### Imputation algorithm 1:

- 01: Input
- 02: m1, m2
- 03: Output
- 04: m1
- 05: Begin
- 06: For i=1 to row no in m2

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07: Evaluate feature index  $F_x = \text{find}(\text{empty}(N_{x_i}))$   
 08:  $F_y = \sim F_x$   
 09: Evaluate  $C_i, c_i$  of the  $m_1$  by K-means  
 10: Index=0  
 11: For  $j=1$  to column no of  $F_x$   
 12: mindist=0  
 13: For  $k=1$  to 6  
 14:  $\text{dist}_k = \|F_{y_i} - C_i\|$   
 15: if  $\text{mindist}_k > \text{dist}_k$  then  
 16:  $\text{mindist}_k = \text{dist}_k$   
 17: Index= $k$ ;  
 18: End if  
 19: Next  $k$   
 20: Impute  $F_x$  with mean values of  $C_{\text{index}}$   
 21: Next  $j$   
 22:  $m_1 = m_1 + (F_{x_i} + F_{y_i})$   
 23: Next  $i$   
 24: End

**Multilayer perception imputation process**

MLP is an artificial neural network in which the hidden layers and neurons in the network topology are determined by a trial and error strategy [12]. An MLP network can undergo a learning process so as to predict MVs. In the MLP imputation process, network consists of one hidden layer and two-layered weight values. The weights of the first layer are related with variables of the input data, and the weights of the second layer are related with the output unit. Features comprising MVs in the training process are used as output while the remaining features are used as input [9]. Training takes place using the complete data, and features including incomplete data are presented to the model for prediction. All features including MVs are presented to the algorithm [13].

**Imputation algorithm 2:**

01: Input  
 02:  $m_1, m_2$   
 03: Output  
 04:  $m_2$   
 05: Begin  
 06: Evaluate feature index  $F_x = \text{find}(\text{empty}(m_2))$   
 07: Set parameters (train\_ratio=70/100, Val\_ratio=15/100, Test\_ratio=15/100)  
 08: For  $i=1$  to col no of  $F_x$   
 09:  $T_i = m_1$ ;

10:  $m_1 = m_1 - T_i$   
 11: Train MLP network by  $T_i, m_1$   
 12:  $In_i = m_2 - F_{x_i}$   
 13:  $F_{x_i} = \text{Test MLP network by } In_i$   
 14:  $m_2 = m_2 + F_{x_i}$   
 15: Next  $i$   
 16: End

**Self-organizing map imputation process**

An SOM is a neural network model comprising an input space and an output layer, and its topology is described on a map. Input values in the training set are described as  $n$ -dimensional  $X = (X_1, X_2, X_3, \dots, X_n)$ . The weight vector of each node in the output layer is  $n$ -dimensional and is described as  $W = (W_1, W_2, W_3, \dots, W_n)$ . The node is described as the Best Matching Node (BMN) to calculate the minimum distance. In the course of MVs prediction, it is observed that the size of the map affects the result. In order to eliminate limit restrictions, a square map that can be created based on the shape is used [14].  $8 \times 8$  SOM network is selected by testing maps of various dimensions ranging between  $4 \times 4$  and  $20 \times 20$ . While dimensions smaller than  $8 \times 8$  do not present the data distribution satisfactorily, larger models are observed to fit the training data.

The cases including MVs are given to the SOM network in turn for prediction. When the value of a feature in an incomplete case is presented to the network for prediction, input values comprising MVs are ignored for the selection of the BMN if they already exist in the case. The steps of the MV imputation algorithm are as follows.

**Imputation algorithm 3:**

01: Input  
 02:  $m_1, m_2$   
 03: Output  
 04:  $m_1$   
 05: Begin  
 06: For  $i=1$  to row no in  $m_2$   
 07: Evaluate feature index  $F_x = \text{find}(\text{empty}(N_{x_i}))$   
 08: Set parameters  
 09: epochs=1000  
 10: Perform  $F_{cn} = \text{SSE}$   
 11:  $\text{mapdim} = 8 \times 8$   
 12: Evaluate  $\text{train}C_i, \text{test}C_i$   
 13: For  $j=1$  to col no of  $\text{test}C_i$   
 14: For all  $w_{ij} \in W$   $d_{ij} = \|N_{x_i} - w_{ij}\|$   
 15: Evaluate weight values of activation group of BMU

- 16:  $W(t+1) = W(t) + \Theta(t)(D_{ij}(t) - W(t))$
- 17: Impute  $N_{x_{ij}}$  with mean values of  $W(t+1)$
- 18: Next  $j$
- 19:  $m1 = m1 + N_{x_i}$
- 20: Next  $i$
- 21: End

**Building the classifiers**

Classification algorithms are one of the key points which have the ability to provide correct information for the evaluation to any algorithm within a machine learning framework. From this point of view, in this study classification algorithms are utilized to judge on how well the resultant dataset is classified. Support Vector Machine (SVM) [15], MLP [16,17], RF [18], and LMT [19] are used to evaluate performances due to successful applications in medical data in recent years. All of the input parameters of classifiers are given as Weka default parameters.

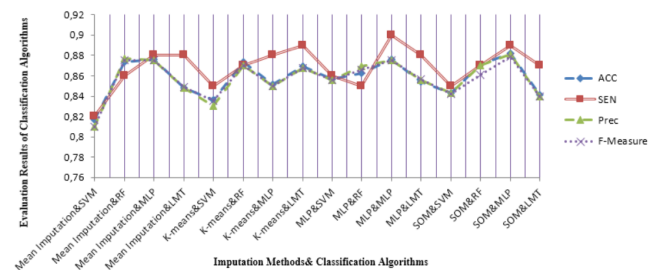
**Results**

The results are evaluated for accuracy (ACC), specificity (Spec), F-measure, and precision (Prec), sensitivity (Sen) in terms of certain metrics. Moreover, the MVs-deleted dataset condition was used as a baseline and the classification performance on datasets obtained as a result of imputation methods is presented in Table 3.

It is significant to evaluate the prediction performance of classifiers for CAD from different perspectives in clinical data analysis. Performance results based on evaluation metrics are presented in Figure 1.

The results indicated 87.58% accuracy on two datasets that are imputed using both the mean method and the MLP imputation method and tested by the MLP classifier. In addition, this dataset has the highest sensitivity, with a value of 0.9, as seen in Table 3. It is observed that the lowest accuracy (81.69%) and the lowest sensitivity (0.82) were obtained in the tests conducted using the SVM classifier.

For the dataset prepared by the SOM imputation method, MLP achieved an accuracy ratio of 88.23%, which was the top value among the imputation-classification method pairs. This combination performed better than machine learning and mean method with regard to both MV imputation and classification. Moreover, other metrics were evaluated for all imputation-classification pairs and SOM imputation method has the best values obtained by MLP, with the accuracy of 88.23%, precision of 0.88, and F-measure of 0.879. Other metric results are presented in Figure 2.

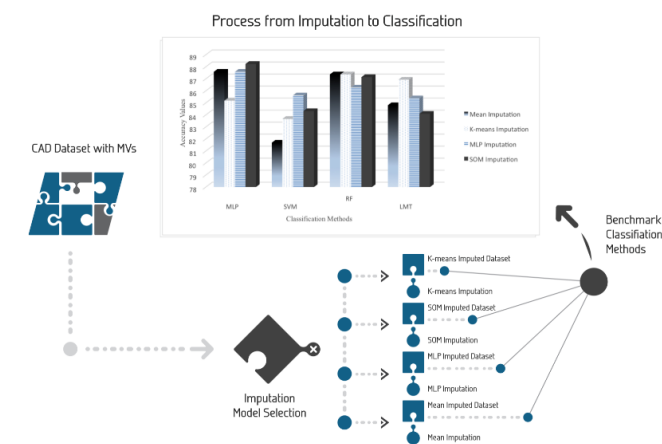


**Figure 2.** ROCs of classification results of mean, K-means, SOM, and MLP imputed datasets in turn with MLP, LMT, SVM, and RF classifiers.

Moreover, according to the average values of the classification results which are presented in Table 3, it is found that the dataset imputed by MLP is more stable than other imputed datasets and increases the performance of all classifiers in the study.

**Table 3.** Classification results from different type of missing value replacement methods.

		ACC	Sen	Spec	F <sub>meas</sub>	Prec
Mean imputation	SVM	81.69	0.82	0.17	0.81	0.81
	RF	86.92	0.86	0.09	0.875	0.877
	MLP	87.58	0.88	0.15	0.875	0.875
	LMT	84.8	0.88	0.23	0.849	0.848
	Average	85.25	0.86	0.16	0.85	0.85
K-means imputation	SVM	83.66	0.85	0.19	0.835	0.83
	RF	87.36	0.87	0.1	0.87	0.87
	MLP	85.18	0.88	0.22	0.85	0.85
	LMT	86.92	0.89	0.19	0.868	0.868
Average	85.78	0.87	0.18	0.86	0.85	
MLP imputation	SVM	85.62	0.86	0.15	0.856	0.856
	RF	86.27	0.85	0.09	0.866	0.869



**Figure 1.** Accuracy results of classification methods for mean, K-means, SOM, and MLP imputed datasets.

For the original part of dataset with 277 samples, LMT achieved an accuracy ratio of 76.17% and the sensitivity value of 0.51, which was the top value for this dataset. This accuracy value remains under 7.49 points obtained by K-means imputation method and tested by the SVM classifier which is the lowest performance between imputations-classifications pairs.

	MLP	87.58	0.9	0.18	0.875	0.875
	LMT	85.38	0.88	0.19	0.857	0.856
Average		86.21	0.87	0.15	0.86	0.86
SOM imputation	SVM	84.31	0.85	0.17	0.842	0.843
	RF	87.14	0.87	0.11	0.861	0.87
	MLP	88.23	0.89	0.1	0.879	0.881
	LMT	84.09	0.87	0.22	0.84	0.84
Average		85.94	0.87	0.15	0.86	0.86
Deleting cases	SVM	74.76	0.51	0.19	0.74	0.73
	RF	75.81	0.5	0.21	0.72	0.72
	MLP	72.92	0.44	0.17	0.73	0.73
	LMT	76.17	0.51	0.19	0.74	0.73
Average		74.92	0.49	0.19	0.73	0.73

## Discussion

Techniques developed for diagnosing CAD should provide considerable diversity in the style of clinical datasets. Sample size should be satisfactory and the connections between features should be determined. In this study, we propose a set of features relating to traditional risk factors such as gender, age, family history concurrently new para-clinical features such as RDW, MCV that were tested and proved to be efficient for identifying cases of CAD are included the model. The aim is to solve the MV problem of which these features store and finally to make a more stable diagnosis using imputed datasets. This study suggests alternative imputation procedures in the case of the lack of laboratory test as well other specific parameters. Thus, a cost-effective technique is constructed that does not need all potential laboratory tests for each suspect individual. It is a comforting situation for the patients on the financial terms.

When the MLP classifier is used on the dataset to which the MLP imputation algorithm is applied, it produces better results than the other methods, with a sensitivity of 0.9 and a specificity value of 0.18. According to the average classification results, the MLP imputation method produces more stable results than all the methods operated with different types of classifiers. The dataset produced by using the mean imputation method, which is widely used by researchers and has been used in this study as a reference, ranks fourth among the algorithm applications.

It has been inferred that MLP in particular is the most effective approach in the MV imputation process for the CAD dataset that includes continuous values when it is particularly tested with machine learning classifiers.

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