



NOVEL MULTIPLE IMPUTATION COMPARISON WITH SIMPLE LINEAR CLASSIFIER, SUPPORT VECTOR MACHINE AND NAÏVE BAYES CLASSIFIER

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ABSTRACT: Incomplete data is a common obstacle to the analysis of data in a variety of fields, ranging from clinical trials to social sciences. Missing values can occur for several different reasons including failure to answer a survey question, dropout, planned missing values, intermittent missed measurements, latent variables, and equipment malfunction. Multiple imputations is one method for handling incomplete data that accounts for the variability of the incomplete data. This procedure does so by filling in plausible values several times to create several complete data sets and then appropriately combining complete data estimates using specific combining rules. We introduced the methodology of multiple imputations in multiple stages and the associated comparison of Simple Linear Classifier, Support Vector Machine and Naïve Bayes Classifiers needed for implementation. We demonstrated via simulations that we have an efficient estimator under the assumption.

Keywords: [Multiple Imputation, Classifier, SVM, Naïve Bayes, Missing Data]

1. INTRODUCTION



Figure 1: Data analysis in Multiple imputation

Possible imputation should give sensible expectations for the missing data, and the inconstancy among them must mirror a fitting level of unsteadiness. Rubin³ recommends that imputations be made through Bayesian disputes: Specify a parametric model for the entire data under MAR, expect a prior appropriation for the dark model parameters, and reenact diverse self-governing draws from the restrictive dispersion of missing esteems given watched data by Bayes hypothesis. Diverse imputation models have been delivered within more expansive and entangled settings. The MATLAB code for REALCOM to extend it to incorporate prior information to allow MNAR imputation, as described below. The REALCOM software uses a joint multivariate normal modelling approach through the Bayesian estimation method Markov Chain Monte Carlo (MCMC).

2. LITERATURE SURVEY

<p>1. Melanie Smuk</p>	<p>2015</p>	<p>Multiple imputations (MI) are a popular tool used to fill in partially observed data with plausible values drawn from an appropriate imputation distribution. Software generally implements MI under the assumption that data are 'missing at random' (MAR) i.e. that the missing mechanism is not dependent on the missing data conditional on the observed data. Broadly there are two ways to frame, and perform sensitivity analyses (SA) to accomplish this: using a pattern mixture model or a selection model. Motivated by a cancer dataset, we develop a novel pattern mixture approach to collecting and incorporating in the analysis prior information elicited from experts. We demonstrated the inferential validity of our approach by simulation.</p>
<p>2. Giuseppe DiCesare</p>	<p>2006</p>	<p>In the special case of Gaussian stochastic processes the problem is simplified since the conditional finite dimensional distributions of the process are multivariate Normal. For more general diffusion processes, including those with jump components, an acceptance-rejection simulation algorithm is introduced which enables one to sample from the exact conditional distribution without appealing to approximate time step methods such as the popular Euler or Milstein schemes. The method is referred to as path wise imputation. Its practical implementation relies only on the basic elements of simulation while its theoretical justification depends on the path wise properties of stochastic processes and in particular Girsanov's theorem.</p>
<p>3. JesperHornblad</p>	<p>2013</p>	<p>In this master thesis, complete case analysis, unconditional mean imputation as well as single and multiple imputation under a fully conditionally specified model were used to impute the diagnosis related group weights and calculate the case mix index. The analysis of their performance showed that all methods produced almost unbiased estimates as</p>

		long as the data was missing completely at random. In contrast, when the missing data mechanism was depending on the value of the diagnosis related group weights, all methods produced biased results. Both, single and multiple imputation noticeably reduced the bias compared to complete case analysis and unconditional mean imputation.
4. Jerome P. Reiter		This article presents an approach for generating multiply-imputed, partitioned synthetic data sets that simultaneously handles the disclosure limitation and missing data. The basic idea is to fill in the missing data first to generate m completed datasets, then replace sensitive or identifying values in each completed dataset with impute values. This article also develops methods for obtaining valid inferences from such multiply-imputed datasets.
5. Matteo Quartagno	2016	In this thesis we propose a Joint Modelling Multiple Imputation (JM-MI) approach to overcome these issues. Motivated by the lack of available software, in the first part of this thesis we develop and describe jomo, a new R package for Multilevel MI. A key feature of jomo compared to other packages for MI, is that it allows for the presence of random, or fixed, study-specific covariance matrices in the imputation model, therefore allowing for heteroscedasticity when imputing.
6. Catherine Welch	2015	This paper propose to adapt, evaluate and implement the two-fold FCS algorithm to impute missing data from large primary care database. To achieve this, first investigate the extent and patterns of missing data in a longitudinal clinical database for health indicators associated with cardiovascular disease risk to determine if the MAR assumption is plausible. Additionally, develop methods to identify and remove outliers, which can potentially bias imputations, from data with repeated measurements before imputation. Adapt and develop the two-fold FCS multiple imputation algorithm to impute missing values in longitudinal clinical data for health indicators associated with cardiovascular disease risk and validate the two-fold FCS algorithm to assess bias and precision through challenging simulation studies.
7. Brian Tinnell Keller	2015	This research serves two purposes: (1) to develop an algorithm in order to implement FCS in the context of a three-level model and (2) to evaluate both imputation methods. The simulation investigated a random intercept model under both 20% and 40% missing data rates. The findings of this thesis suggest

		that the estimates for both JM and FCS were largely unbiased, gave good coverage, and produced similar results.
8. Yang Yuan	2014	The first part placed a particular emphasis on the so called missing at random (MAR) assumption, but focuses the bulk of attention on multiple imputation techniques. The main aim of this part is to investigate various modelling techniques using application studies, and to specify the most appropriate techniques as well as gain insight into the appropriateness of these techniques for handling incomplete data analysis. This thesis first deals with the problem of missing covariate values to estimate regression parameters under a monotone missing covariate pattern. The study is devoted to a comparison of different imputation techniques, namely markov chain monte carlo (MCMC), regression, propensity score (PS) and last observation carried forward (LOCF). The results from the application study revealed that we have universally best methods to deal with missing covariates when the missing data pattern is monotone.

3. PROPOSED WORK

3.1 NOVEL MULTIPLE IMPUTATION COMPARISON WITH SIMPLE LINEAR CLASSIFIER, SUPPORT VECTOR MACHINE AND NAÏVE BAYES CLASSIFIER

The major objective of this paper is to implement and compare the proposed framework with three classification Simple Linear Classifier , Support vector Machine, Naïve Bayes Classifier to build up an automated decision support framework for Multiple Imputation practice. The design was to decide an ideal classification mechanism for Multiple Imputation plans with high diagnostic accuracy. Distinctive classification algorithms were tried and benchmarked for their performance. The performance of the classification algorithms is illustrated on benchmark datasets. A linear classifier achieves this by making a classification decision based on the value of a linear combination of the characteristics. A question's characteristics are also known as feature values and are typically displayed to the machine in a vector called a feature vector. we demonstrate that the two learning methods Naive Bayes and Rocchio are instances of linear classifiers, the perhaps most important gathering of content classifiers, and contrast them with nonlinear classifiers. To streamline the talk, we will just consider two-class classifiers in this segment and characterize a linear classifier as a two-class classifier that decides class enrollment by comparing a linear combination of the features to a limit.

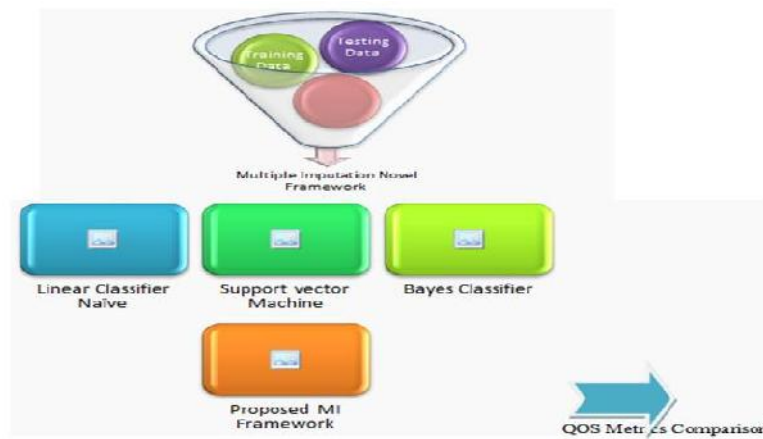


Figure 2: Proposed work Overflow

SVMs were first proposed by Vapnik in the 1960s for classification and have as of late turned into an area of exceptional research inferable from improvements in the methods and hypothesis combined with augmentations to relapse and thickness estimation. SVMs arose from statistical learning hypothesis; the aim being to take care of just the problem of enthusiasm without taking care of a more troublesome problem as an intermediate advance. SVMs are based on the structural hazard minimization principle, firmly related to regularization hypothesis. This principle incorporates capacity control to counteract overfitting and consequently is a partial answer for the bias-variance trade-off dilemma.

4. EXPERIMENTAL RESULTS

Classification is a data mining process that appoints things in an accumulation to target classifications or classes. The objective of classification is to foresee an objective class for each case in the dataset precisely. Table 1 represented into comparison with proposed overall metric values and Figure 3 displayed to comparison values diagram. Classification ability relies upon the kinds of algorithms and the attributes of the data, for example, the level of imbalance, number of highlights, number of instances, and number of class composes. Besides, while missing values are dealt with by a specific imputation method, the classification algorithm is additionally influenced by the imputation method. In this manner, each extraordinary imputation method/classifier combine brings about an alternate execution, regardless of whether they treat similar data with the same missing values. Figure 4 represented into comparison of mean metrics using values.

	Linear Classifier	Support Vector Machine	Naïve Bayes Classifier	Proposed
PCC	-0.1	-0.07	0.06	0.1
Mean Abs Sqr	0.05	0.08	0.09	0.1
RM Sqr Error	25	45	65	85
Precision	0.01	0.05	0.02	0.09
Recall	7	35	42	89
F-Score	75	50	280	390

Table 1: Comparison of proposed overall metrics

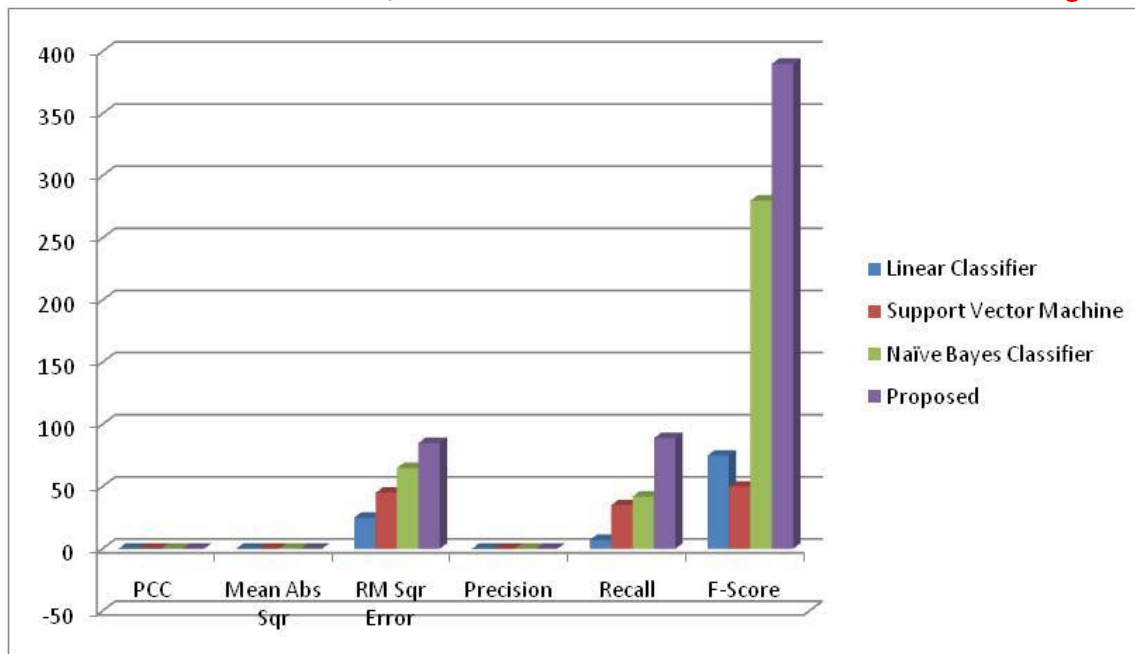


Figure 3: Proposed Overall Metrics Values

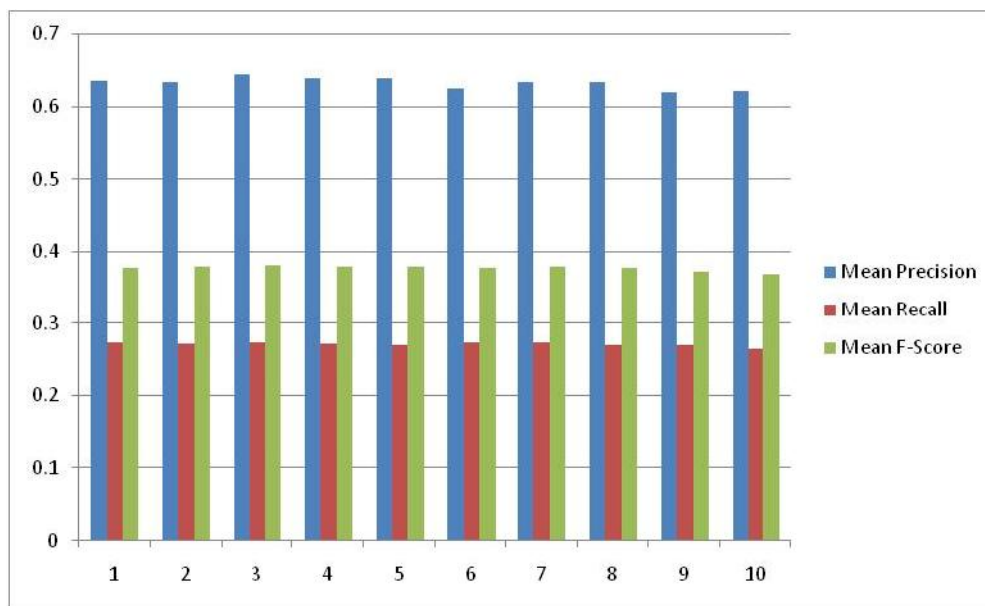


Figure 4: Comparison of Mean metrics using 10 Runs values

CONCLUSION

We presented the Novel methodology of multiple imputations in multiple stages and the related correlation of various classifiers required for implementation. We showed by means of simulations that we have an efficient estimator under the suspicion. We demonstrated that the method functions admirably as far as low percent bias low mean square error and great coverage. The simulations in this article investigated the effect of the request and number of imputations under a predetermined arrangement of conditions, most quite, ignorability. The outcomes demonstrate that neither the request, nor the quantity of imputations have huge effect on the bias, mean square error, or coverage, under this arrangement of conditions. this work gives a pattern framework to more complex situations and more complex suppositions forced on the missing values and classification of missing data.

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