# Multiple Imputation for Missing Data in Repeated Measurements Using MCMC and Copulas

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Abstract— This paper presents two imputation methods: Markov Chain Monte Carlo (MCMC) and Copulas to handle missing data in repeated measurements. Simulation studies were performed using the Monte Carlo technique to generate datasets in different situations. Each subject unit in each dataset was measured on three occasions under the following conditions: 1. data had a multivariate normal distribution under two types of correlation structures: Compound Symmetry (CS) and Autoregressive (AR (1)), 2. the correlation among repeated observations under each subject was determined at low level ( $\rho$  =0.3), middle level ( $\rho$  =0.5), and high level ( $\rho$ =0.7), 3. sample sizes consisted of 30, 70, and 100 subject units, and 4. data were assigned missing at random (MAR) at the last occasion of measurement with missing rate of 5%, 10%, 20% and 30%, respectively. All possible combinations of these conditions gave rise a total of 72 different situations. Each defined situation was repeated 1,000 times by SAS programming and each missing value was replaced with a set of five plausible values that represent the uncertainty about the right value to impute under the MCMC method. The performance of each imputation method was evaluated using mean square error (MSE). The lower MSE would indicate the more effective method. The results from the simulation studies showed that the Copulas method was superior effective than other methods in all situations. The MCMC method was more effective than the simple mean imputation method when the correlation structure was AR1. For application, both imputation methods were applied with two datasets in practices: 1) waist circumference data on healthy project and 2) monthly rainfall data. The results also confirmed that the Copulas was the most effective method which was consistent with the simulation studies.

*Index Terms*— Markov Chain Monte Carlo, Copulas, missing at random, repeated measuremens.

## I. INTRODUCTION

IN repeated measures data analysis, it often faces with the problem of incomplete data when a subject has one or more missing values during the subsequent waves of data collection. The occurrence of missing values may due to any causes related to a study unit such as nonresponse,

refusal, drop out, lost of follow up, illness or death. In the

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meantime, missing values can be generated from human errors on either forget to ask a question or forget to record the answer. The patterns of missing data may be one of these special patterns: 1) univariate missing data, 2) unit nonresponse, or 3) monotone missing data. Firstly, the "univariate" missing data are restricted to missing values on a single variable while the other variables are fully recorded. Secondly, the "unit nonresponse" missing data have missing values on a block of variables for the same set of cases, and the rest of variables are all complete. Thirdly, the "monotone" missing data have the pattern of missing values after all variables are arranged such that  $X_1, X_2, \ldots, X_j$ :  $j = 1, 2, \ldots, k - 1; k > 1$ , then the variable  $X_j$  will be

observed whenever the variable  $X_{i+1}$  is observed [8], [12].

To handle problems of missing data, one simplest approach is to focus on a complete-case analysis, but its disadvantage is the decreasing on statistical power from the smaller sample size [9], [10]. Another approach for analyzing incomplete data is using imputation methods to impute the best estimate of a missing value of the variable [7], [12]. Imputation methods base on three types of missingness as follows: 1) missing completely at random (MCAR, if the missingness is independent of both unobserved and observed data), 2) missing at random (MAR, if the missingness depends on observed data, but not on unobserved data), and 3) missing not at random (MNAR, if the missingness depends on unobserved data) [7], [13]. Multiple imputation (MI), under imputation approach, is proposed by Rubin [2] to analyze incomplete data under MAR mechanism. The idea of MI procedure is to replace each missing value with a set of M possible values. These values are drawn from the distribution of the study data under the uncertainty about the right value to impute. Then, analyze the imputed data sets from standard procedure for complete data and combine the results from these analyses [1], [13]. According to the MI concepts, this study aims to 1) present the possibilities of using two imputation methods: Markov Chain Monte Carlo (MCMC) and Copulas to impute the best estimates of missing values under MAR mechanism in repeated measurements, and 2) compare the performance of the MCMC and Copulas methods in estimating the missing values.

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## II. METHODS

#### A. Generating Datasets with Missing Values

A number of complete datasets were generated to have data on all three repeated measurements. Let  $Y_1, Y_2$ , and

 $Y_3$  be three continuous variables for three measurements on each subject at time point 1, 2, and 3, respectively. Under the assumption of a multivariate normal distribution with zero mean vector and variance-covariance matrix  $\sum_{n=1}^{\infty} \frac{1}{n}$ ,

$$\Sigma = \begin{bmatrix} \boldsymbol{\sigma}_{11} & \boldsymbol{\sigma}_{12} & \boldsymbol{\sigma}_{13} \\ \boldsymbol{\sigma}_{21} & \boldsymbol{\sigma}_{22} & \boldsymbol{\sigma}_{23} \\ \boldsymbol{\sigma}_{31} & \boldsymbol{\sigma}_{32} & \boldsymbol{\sigma}_{33} \end{bmatrix},$$

 $Y_1, Y_2$ , and  $Y_3$  are correlated under two types of

correlation structures, including compound symmetry (CS):

$$R_{cs} = \begin{vmatrix} 1 & \rho & \rho \\ \rho & 1 & \rho \\ \rho & \rho & 1 \end{vmatrix}$$

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and first order autoregressive (AR(1)):

$$R_{AR(1)} = \begin{bmatrix} 1 & \rho & \rho^{2} \\ \rho & 1 & \rho \\ \rho^{2} & \rho & 1 \end{bmatrix},$$
  
where  $\rho_{ij} = \frac{\sigma_{ij}}{\sqrt{\sigma_{ij}}\sqrt{\sigma_{ij}}}; i, j = 1, 2, 3.$ 

The correlation among repeated observations under each subject was determined at low level ( $\rho = 0.3$ ), middle level ( $\rho = 0.5$ ), and high level ( $\rho = 0.7$ ). Sample sizes considered were 30, 70, and 100 subject units. Additionally, data were assumed MAR at the last occasion of measurement,  $Y_3$ , with missing rate of 5%, 10%, 20% and 30%, respectively. All possible combinations of these conditions gave rise a total of  $2 \times 3 \times 3 \times 3 = 72$  different situations for 72 different datasets.

#### B. Imputation Methods

In each dataset, a simple imputation method was used to replace the missing value with a single value of the variable's mean of the complete cases. This single imputation did not reflect the uncertainty about the prediction of the unknown missing values. Then, two advanced imputation methods, MCMC and Copulas, were used to estimate the missing value under MAR mechanism in repeated measures.

## C. MCMC

MCMC is a numerical method for generating pseudorandom drawn from probability distributions via Markov Chains. A Markov chain is a sequence of random variables  $X_0, X_1, X_2, X_3, \dots, X_i$  in which the distribution of each element given all previous ones depends only on the most recent value, i.e. for all i,

$$P[X_{i} < x | X_{0} = x_{0}, X_{1} = x_{1}, \dots, X_{i-1} = x_{i-1}]$$

 $= P[X_{i} < x | X_{i-1} = x_{i-1}].$ 

A chain of random variables must be created long enough to approach to the approximate stationary distribution. By repeated simulation steps of the chain, one simulates draws samples from the stationary distribution [1], [7].

Reference [7] suggested using MCMC to impute missing values by assuming: 1) data is from a multivariate normal distribution, 2) data are MCAR or MAR, and 3) the missing data pattern can be either monotone or arbitrary. To begin the MCMC process, a vector of means ( $\mu$ ) and a variance-

covariance matrix  $(\Sigma)$  from the complete data (or available

cases) are computed and used as the initial estimates for the Expectation-Maximization (EM) algorithm. The resulting EM estimates are used to be the starting values to estimate the parameters of the prior distributions for means and variances of the multivariate normal distribution with informative prior. In MCMC, the imputation step (I-step) simulates values of missing items by randomly selecting a value from the conditional distribution of missing values,  $Y_{i(miss)}$ , given the observed values,  $Y_{i(obs)}$ . Next, the posterior step (P-step) updates the posterior distribution of the mean and covariance parameters (e.g., normal distribution for the means and inverted Wishart distribution for covariance matrix [3]). Then, the vector of means and covariance matrix are simulated from the posterior distribution based on the updated parameters. The new estimates will be used in the I-step. Both the I-step and the P-step are iterated until the mean vector and covariance matrix are unchanged. The imputation from the final iteration will be used to form a complete data set.

This study applied the MCMC method with the predetermined 72 different situations of repeated measurements. Each defined situation was repeated 1,000 times by using SAS procedures and each missing value was replaced with a set of five plausible values from posterior predictive distribution that represent the uncertainty about the right value to impute. Finally, the results from the five plausible values of each missing value were combined and used their average value to be the best estimates of each missing value.

## D. Copulas

In general, the imputation of missing value is based on the conditional distribution that we need to know the joint distribution of repeated measures those are prior to the observation with missing value. Let observation at the  $k^{th}$ occasion of measurement have missing value. The idea of using Copulas is to create a joint distribution from marginal distribution of the 1<sup>st</sup>, 2<sup>nd</sup>, 3<sup>rd</sup>,..., (k-1)<sup>th</sup> occasion of measurement. Then, we can find a conditional distribution of the k<sup>th</sup> measurement given the 1<sup>st</sup>, 2<sup>nd</sup>, 3<sup>rd</sup>,..., and (k-1)<sup>th</sup> measurement [1], [5], [6]. The Copulas is a function that links univariate marginal distributions to their joint multivariate distribution function as the following equation:

 $F(x_{1}, x_{2}, \dots, x_{k}) = C(F_{1}(x_{1}), F_{2}(x_{2}), \dots, F_{k}(x_{k}))$ (1)

where

F is the distribution function on  $\Re^k$ 

 $F_1, F_2, \ldots, F_n$  are univariate marginal distributions,

 $C(F_1(x_1), F_2(x_2), \dots, F_n(x_n))$  is a multivariate

distribution function with marginals  $F_1, F_2, \ldots, F_n$ , and C is copula function.

If one knows the joint distribution of  $x = (X_1, X_2, \dots, X_k)$ , a vector of random variables, then one can impute the missing value from a conditional distribution of  $X_k$ , a vector with missing values, given  $H = (X_1, X_2, \dots, X_{k-1})$  which is a vector with complete data in history.

In this study, we applied the Gaussian copula in which the k-variate Gaussian copula with k Gaussian marginals corresponds to the k-variate Gaussian distribution. That is the multivariate normal distribution has normal marginal distributions and Gaussian copula dependence. Gaussian copula handles the dependence among univariate marginal distributions via the correlation matrix R of pairwise dependencies between variables. The multivariate Gaussian copula is defined as:

$$C_k(u_1, u_2, ..., u_k; R) = \Phi_k(\Phi_1^{-1}(u_1), \Phi_1^{-1}(u_2), ..., \Phi_1^{-1}(u_k))$$

where  $u_j \in (0,1)$ , j = 1, 2,...,k;  $\Phi_k$  is the standard k variate normal distribution function with the correlation matrix R. For modeling the repeated measurements using The multivariate Gaussian copula, let  $X_j$  have a continuous distribution function  $F_j$ , j = 1,2,..., k, and let  $Y_j$  be its normalizing transformation as  $Y_j = \Phi_1^{-1}[F_j(X_j)]$ , where  $\Phi_1^{-1}$  is the inverse of the standard univariate Gaussian distribution function. Then, the joint multivariate distribution function is:

$$F(y_{1}, y_{2}, ..., y_{k}; R) = C_{k}(u_{1}, u_{2}, ..., u_{k}; R) = \Phi_{k}[\Phi_{1}^{-1}(u_{1}), \Phi_{1}^{-1}(u_{2}), ..., \Phi_{1}^{-1}(u_{k}); R].$$
(2)

Then, we can find the MLE of missing value  $y_k$  from the conditional distribution of  $f(y_k | H, R)$ . Lastly, the missing value ( $y_k$ ) can be replaced by:

$$\hat{y}_{k}^{CS} = \frac{\rho}{1 + (k-2)\rho} \sum_{j=1}^{k-1} y_{j}; \text{ if } R \text{ is under CS structure,}$$

 $\text{ and } \ \hat{y}_{k}^{\text{AR}} = \rho \frac{S_{k}}{S_{k-1}} \left( y_{k-1} - \overline{Y}_{k-1} \right) + \ \overline{Y}_{k} \, ; \ \text{if } \text{ R} \ \text{is under}$ 

AR1 structure, where  $S_{k-1}$  and  $S_k$  are standard deviation at the (k-1)<sup>th</sup> and the k<sup>th</sup> occasion of measurement [5], [6].

# E. Comparison of Imputation Methods

Once the MCMC method has been implemented, it is necessary to check the convergence of the simulated sequence of random variables to the stationary distribution. However, it is not too difficult to apply the MCMC method for multiple imputations of missing values because it has commands available on some statistical packages such as PROC MI and PROC MINIMIZE in SAS program [1], [13]. In opposite, the copulas method is easy for computation but it is not easy to replace the values of missing data when there are lots of missing items. In addition, the data on repeated measurements those are prior to the observation of missing value must be complete. It also needs to check whether the correlation structure among repeated observations is under CS or AR(1). The performance of each imputation method can be evaluated from its value of mean square error (MSE). The lower MSE would indicate the more effective method [10].

## III. RESULTS

To meet the objectives of study, we compared among three imputation methods: simple mean, MCMC and Copulas for the best estimates of missing values under MAR mechanism in repeated measurements both in simulation and application of two real datasets.

## A. Simulation Study

Table 1 and Table 2 showed the estimated MSE from simulation studies in 72 situations which are under different types of correlation structure, levels of correlation among repeated measurements, missing rates (%) at last measurement, and sample sizes. Results indicated that the Copulas method was the most effective in all situations. The MCMC method was more effective than the simple mean imputation method when the correlation structure was AR(1).

# B. Application of Two Datasetss

We applied the MCMC and the Copulas algorithms for imputation of missing values in two real datasets with different correlation structure among repeated observations.

## Dataset 1: The waist circumference data

The first dataset was obtained from a study on "healthy project" of the "Nopparat Rajathanee" hospital in Bangkok where the outcome of interest was waist circumference of each individual. The waist circumference data, including weight and height, were collected from participants on a monthly basis between November 2008 and February 2009. At least 105 participants have waist circumference values more than 80 and 90 centimeters and 44 participants were completely followed up for four months. The correlation among repeated observations of waist circumference under each subject was assigned at high level ( $\rho = 0.91625 \approx 0.9$ ) from the assuming CS correlation structure as data evidence-based shown in the below:

$R_{cs} =$	1	0.93662	0.92046	0.85249	
	0.93662	1	0.94599	0.83225	
	0.92046	0.94599	1	0.91625	
	0.85249	0.83225	0.91625	1	

The data on the last occasion of measurements were assigned missing at random with missing rate of 5%, 10%, 20% and 30%, respectively. Next, we tested for the normal distribution of the waist circumference data containing missing values, checked for convergence of mean vector and covariance matrix under the MCMC process for MI, and compared the performance of three imputation methods: simple mean, MCMC, and Copulas. The result of study showed that the MSE of the Copulas method is smallest at all level of missing rate (Fig. 1(a).) when compares with those of other imputation methods. For this dataset, the Copulas shows the most effective method for data imputation under CS correlation structure.

#### Dataset 2: The monthly rainfall data

The second dataset was from routine work of Thai Meteorological Department (TMD) where the outcomes of interest are the average of monthly rainfall in the northern region of Thailand. The data were collected from 28 local weather stations from June to August, 2009. The correlation among repeated observations of the average of monthly rainfall at each weather station was assigned at middle level ( $\rho = 0.452 \approx 0.5$ ) from the assuming AR(1) correlation structure as data evidence-based shown in the below:

$$R_{AR(1)} = \begin{vmatrix} 1 & 0.3566 & 0.189 \\ 0.3566 & 1 & 0.452 \\ 0.189 & 0.452 & 1 \end{vmatrix}$$

The data on the average rain volume in August 2009 were assigned missing at random with missing rate of 5%, 10%, 20% and 30%, respectively. Next, we tested for the normal distribution of the average of monthly rainfall data which included missing values, checked for convergence of mean vector and covariance matrix under the MCMC process for MI, and compared the performance of the three imputation methods: simple mean, MCMC, and Copulas. The result of study showed that the MSE of the Copulas method is smallest at all level of missing rates (Fig. 1(b).) when compares with other imputation methods. For this dataset, the Copulas also shows the most effective method for data imputation under AR(1) structure.

### IV. DISCUSSION

Although the results from simulation studies indicated that the Copulas method was superior to the MCMC and the simple mean imputation methods, its performance depended on these factors, including missing rate (%) and level of correlation among repeated observations. If the sample size was fixed and the missing rate increased from 20% to 30%, there were much more difference in the values of MSE obtained from either the Copulas or the MCMC method. Except for the fixed sample size 70 under the CS correlation structure of repeated observations, the estimated MSE from

ISBN: 978-988-19251-9-0 ISSN: 2078-0958 (Print); ISSN: 2078-0966 (Online) both the Copulas and the MCMC method under missing rate 20% or 30% were closer in values. That is the performance of both imputation methods are almost equivalence when the level of missing rate was increased and the sample size is 70 or larger. Moreover, we found that the MSE values from all imputation methods of the simple mean, the MCMC and the Copulas would decrease when the level of correlation among repeated observations was increased under the condition of fixed sample size and fixed level of missing rate. In sum, the performance of the simple mean and the Copulas methods were higher than the MCMC when the level of correlation among repeated observations was increased under the CS structure. The MCMC and the Copulas gave better performance than the simple mean method under AR(1) structure when the level of correlation among repeated observations was increased.

#### V. CONCLUSION

The results from the simulation studies showed that the Copulas method was the most effective in all situations. The MCMC method was more effective than the simple mean imputation method when the correlation structure was under AR(1). For application, both imputation methods were applied with two datasets in practices: 1) waist circumference data on healthy project and 2) monthly rainfall data.

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Table 1

Comparison of MSE among three imputation methods (simple mean, MCMC, and Copulas) from the simulation study under CS correlation structure, level of correlation among repeated measurement ( $\rho$ ), missing rate of last measurement (%) and sample size

Sample size	missing rate (%)	MSE									
		$\rho = 0.3$				$\rho = 0.5$			$\rho = 0.7$		
		Mean	MCMC	Copulas	Mean	MCMC	Copulas	Mean	MCMC	Copulas	
30	5	1.0250	1.0171	0.8985	0.6712	0.7434	0.6539	0.5321	0.6773	0.4351	
	10	1.0240	1.0229	0.8536	0.7347	0.7782	0.6610	0.5431	0.5790	0.4436	
	20	1.0421	1.0440	0.8595	0.7221	0.8705	0.6724	0.5512	0.5494	0.4492	
	30	1.0500	1.1163	0.8561	0.7591	0.8942	0.6493	0.5578	0.5673	0.4343	
70	5	1.0399	1.0536	0.8298	0.7386	0.8930	0.6587	0.4631	0.5678	0.4399	
	10	1.0449	1.0620	0.8417	0.7783	0.8977	0.6583	0.4849	0.5679	0.4403	
	20	1.0872	1.0643	0.8484	0.7646	0.8427	0.6612	0.4690	0.5355	0.4418	
	30	1.0295	1.0137	0.8518	0.7581	0.7869	0.6830	0.4920	0.5008	0.4642	
100	5	1.0399	1.0079	0.8636	0.8515	0.8882	0.6796	0.5703	0.5663	0.4516	
	10	1.0210	1.0620	0.8648	0.8223	0.8270	0.6760	0.5349	0.5273	0.4494	
	20	1.0584	1.1036	0.8668	0.7874	0.8703	0.6722	0.4923	0.6480	0.4473	
	30	1.0558	1.0751	0.8638	0.7770	0.8383	0.6717	0.4806	0.5326	0.4473	

Table 2

Comparison of MSE among three imputation methods (simple mean, MCMC, and Copulas) from the simulation study under AR(1) correlation structure, level of correlation among repeated measurement ( $\rho$ ), missing rate of last measurement (%) and sample size

Sample size	missing rate (%)	MSE								
		$\rho = 0.3$			ρ = 0.5			$\rho = 0.7$		
		Mean	MCMC	Copulas	Mean	MCMC	Copulas	Mean	MCMC	Copulas
30	5	1.2283	1.1165	0.9350	1.2283	0.9196	0.7777	0.6394	0.6253	0.5367
	10	1.4438	1.0514	0.9220	1.2251	0.8479	0.7643	0.6417	0.5736	0.5268
	20	1.2449	1.1752	0.9342	0.9884	0.9110	0.7776	0.6525	0.6451	0.5393
	30	1.2571	1.2290	0.9667	0.9974	0.9448	0.8049	0.6972	0.6860	0.5600
70	5	1.2321	0.9474	0.8125	0.9805	0.8544	0.7651	0.6983	0.6149	0.5698
	10	1.1231	0.8816	0.8108	0.9762	0.8816	0.7594	0.6471	0.5960	0.5682
	20	1.3218	0.9089	0.8091	0.9790	0.9089	0.7638	0.6571	0.6175	0.5685
	30	1.1973	0.9196	0.8065	0.9812	0.9196	0.7645	0.6982	0.6671	0.5687
100	5	1.2443	1.1152	0.9250	0.9982	0.9187	0.7631	0.6831	0.6243	0.5655
	10	1.2579	1.1126	0.9314	0.9976	0.9176	0.7685	0.6580	0.6242	0.5682
	20	1.2680	1.2100	0.9350	0.9821	0.8942	0.7718	0.6972	0.6757	0.5712
	30	1.2633	1.1391	0.9342	0.9781	0.8946	0.7715	0.7123	0.6369	0.5722



Fig. 1(a). MSE from three imputation methods (simple mean, MCMC, and Copulas) from the waist circumference data. Fig. 1(b). MSE from three imputation methods (simple mean, MCMC, and Copulas) from the monthly rainfall data.