

# New Framework of Location Model for Mixed Variables Classification in the Presence of Outliers

Kartini Kasim<sup>1,2,[\\*](#page-0-0)</sup>, Hashibah Hamid<sup>1</sup>, Ayu Abdul-Rahman<sup>1</sup>

<sup>1</sup> Department of Mathematics, School of Quantitative Sciences, UUM College of Arts and Sciences, Universiti Utara Malaysia, 06010 UUM Sintok, Kedah, Malaysia

<sup>2</sup> School of Mathematical Sciences, College of Computing, Informatics and Mathematics, Universiti Teknologi MARA Kedah Branch, Sungai Petani Campus, 08400 Sungai Petani, Kedah, Malaysia



#### **1. Introduction**

Real-world applications in medical [1-4], engineering [5-8], psychology [9,10] and agriculture [11,12] serve as instances of classification studies dealing with a mixed of continuous and categorical variables. Numerous statistical techniques are employed specifically for the classification task containing mixed variables. One of the technique is the Location Model (LM), a parametric approach that effectively classifies objects consisting of mixed variable types concurrently [13-15]. The LM has demonstrated considerable success in both methodological and model development, as well as its practical application in solving the problem of classification involving a mixture of variables [16-18].

On the other hand, Olkin and Tate [19] was initially introduced the LM by deriving the mixed variables distribution. Later, Chang and Afifi [20] proposed a parametric classification approach that specifically dealt with mixed variables problems. The authors implemented the LM in the

<span id="page-0-0"></span><sup>\*</sup> *Corresponding author.*

*E-mail address: ms.kartini@gmail.com*

classification model for one binary and one continuous variable. In the approach, categorical variables are treated as binary variables with values of zero or one. Consequently, Krzanowski [21] extended the model for more than two variables, followed by another generalization of mixed variables (categorical and continuous) of the two groups' problems [14-22].

Previous research has extensively explored classification procedures utilizing the LM. Table 1 provides examples of such studies, which delve into challenges including high dimensionality, variable selection, and handling a substantial number of binary variables.

#### **Table 1**





All the studies demonstrated great achievement in method and model expansion, as well as successfully solving classification problems regarding mixed variables. However, despite these successes, the classification procedure based on LM only performs well under non-contaminated data scenarios, i.e., free from outliers [26-27]. The definition of outlier has been defined by various perspectives depending on the applications. As exhibited in Table 2, the definitions highlight deviations of data from most of the observed data.

#### **Table 2**

Definition of outlier



The outliers will disrupt the parametric nature of LM, as it relies on the assumption of normality, i.e.,  $(y \sim MVN(\mu_{im}, \Sigma)$ . The presence of outliers in the dataset violate this assumption, affecting key parameters such as the sample mean and covariance. Under this circumstance, the outlier affects the parameters, i.e., sample mean  $(\mu_{im})$  and covariance  $(\Sigma)$ . Consequently, the sample mean may fail to

represent the true center of the distribution, and the covariance may inaccurately capture data variability. This misrepresentation can lead to underestimated or overestimated parameter values, ultimately disrupting the normality assumption and increasing the misclassification rate, resulting in poor performance of the LM [34-36]. To address this gap, the objective of this study is to propose a new framework, the Robust Location Model (RLM<sub>med</sub>), which integrates a median estimator and a robust covariance matrix to replace the classical mean and covariance in the LM, thereby mitigating the impact of outliers on parameter estimation and enhancing the model performance.

# **2. The Classical Location Model**

In the classical LM, there are two groups denoted as  $\pi_i$  ( $i = 1, 2$ ) for Groups 1 and 2, respectively, with  $n_i$  represents the group sample size. The structure of the LM is  $s = 2<sup>b</sup>$  in which *s* represent the number of cells and *b* is the number of binary variables. In the structure, the binary variables drop as cells, and each cell contains only *c* continuous data. Figure 1 illustrates the structure of the LM, with x and y representing binary and continuous variables, respectively. Let  $b = 2$ , which then yields  $s = 4$  per group.



**Fig. 1.** The Structure of the LM when  $b = 2$ 

The continuous data in each cell follows normal distributions with different means and equal covariance within the groups. The probability of obtaining an object in the cell m of  $\pi_i$  is  $p_{i_m}$ . All objects can be observed as a vector in the form of  $z' = (x^T, y^T)$ , where  $x^T = (x_1, x_2, ..., x_b)$  is the vector of *b* binary variables, while  $y^T = (y_1, y_2, ..., y_c)$  is the vector of  $c$  continuous variables. Hence, the conditional distribution of vector y given x for  $\pi_i$  is  $(y|x = m \sim MVN(\mu_{im}, \Sigma))$ . Thus, the optimal rule of the classical LM is written as Eq. (1). The future objects,  $y$  are classified into  $\pi_1$  if

$$
\left(\mu_{1m} - \mu_{2m}\right)^T \sum \left(\nu_r - \frac{1}{2} \left(\mu_{1m} + \mu_{2m}\right)\right) \ge \log \left(\frac{p_{2m}}{p_{1m}}\right),\tag{1}
$$

otherwise, y will be classified into  $\pi_2$ .

In constructing the LM, the presented outliers will distort the values of parameters [35-37]. In this case, the parameters are underestimated (or overestimated), yielding inaccurate classification rules and leading to a high misclassification rate [30,34-36]. Therefore, outliers in the dataset must be tackled prior to constructing the LM to obtain accurate and optimal results.

# **3. RLMmed**

*3.1 Median and Robust Covariance as Alternative to Classical Estimator*

One way to alleviate the effect of outliers is by estimating the parameters using robust estimators [38-42]. Robust estimators are used in place of classical estimators since the former is able to reduce the effect of outliers [43-46]. Furthermore, implementing robust estimators in the model helps minimize the misclassification rate [47-49]. On the other hand, Bickel [50] highlighted that medians are alternative location estimators to the classical mean. Its robustness is evident as it remains unaffected by outliers in up to 50% of the data [51-52]. Alternatively, Croux and Dehon [53] asserted that using a robust covariance matrix in classification analysis can effectively reduce misclassification rates for low and high-contaminated data. Therefore, employing a paired robust covariance and location estimator is recommended to solve the sensitivity toward outliers [54-55]. For this purpose, the mean vectors and covariance matrix in the classical LM are substituted with the median and robust covariance matrix.

# *3.2 New Framework of LM*

In developing the new framework of LM, the outliers are managed prior to its construction. The entire procedure is structured as follows.

# *3.2.1* Phase I: Handling Outliers in the LM

For the first phase, the raw data are split into testing and training sets. In the testing dataset, an object, *k* where  $k = 1, 2, ..., n$  is omitted sequentially, while the remaining object,  $n - k$  is treated as the training set. The training set is used to estimate the median and the robust covariance matrix. The median estimator are computed using the following Eq. (2).

$$
\tilde{y} = \begin{cases} y_{\frac{n+1}{2}} & \text{if } n \text{ is odd} \\ \frac{1}{2} \left( y_{\left(\frac{n}{2}\right)} + y_{\left(\frac{n}{2}\right)+1} \right) & \text{if } n \text{ is even} \end{cases}
$$
 (2)

In this study, the robust covariance matrix is estimated based on the product of Spearman's correlation ( *ρ<sup>s</sup>* ) and the median absolute deviation about the median (MAD*n*). The computation of  $\rho_s$  is computed following Eq. (3).

$$
\rho_s = 1 - \frac{6\sum d^2}{n(n^2 - 1)},
$$
\n(3)

where  $d^2$  is the square of the difference between the ranks of the  $r^{\dot{n}}$  objects of  $y_{r1}$  and  $y_{r2}$ . Meanwhile, the MAD*n* is computed as in Eq. (4).

$$
\text{MAD}n = h \text{ med } |y_r - \text{med}(y_r)|,\tag{4}
$$

where *h* is a constant set at 1.4826 to make the estimator consistent and unbiased for the estimation of standard deviation under a normal distribution [56-57]. The robust covariance matrix  $(S<sub>R</sub>)$  is estimated by multiplying the  $\rho_s$  and MADn as indicated in Eq. (5).

$$
S_R = \rho_s \left( \text{MAD} n_1 \right) \left( \text{MAD} n_2 \right). \tag{5}
$$

## *3.2.3* Phase II: Construction of new LM

In constructing the new LM, the median estimator and the robust covariance matrix are fused into the LM, in which the fusing estimator becomes a new approach. This new approach is applied to the LM, creating a new LM,  $RLM_{med}$ . The  $RLM_{med}$  is computed as indicated in Eq. (6).

$$
\left(\tilde{\mathbf{y}}_{1m} - \tilde{\mathbf{y}}_{2m}\right)^{T} \sum_{s_{R}}^{-1} \left[ y - \frac{1}{2} \left(\tilde{\mathbf{y}}_{1m} + \tilde{\mathbf{y}}_{2m}\right) \right] \ge \log \left(\frac{p_{2m}}{p_{1m}}\right). \tag{6}
$$

## *3.2.3* Phase III: Model validation

In this phase, the newly constructed model,  $RLM_{med}$ , is compared to other classification methods for validation purposes. The performance of  $RLM_{med}$  is measured using the error rate through the leave-one-out method, where the method with the lowest error is considered the best. The RLM $_{med}$ will validated using a real dataset, i.e., an echocardiogram dataset. The echocardiogram dataset consists of 64 patients, of which 41 have not survived less than a year  $(n_1)$ , and the remaining being

23  $(n_2)$ . This data contains nine continuous variables with two binary variables.

## **4. Conclusion**

From this study, the findings are expected to be obtained:

- i. An estimator that is able to handle outlier's problem, which became a new approach in the LM;
- ii. A new LM that can be used as an alternative to other mixed variables classification models. This new LM is expected to perform better than the classical LM.

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