Comparison of Sampling Size Estimation Techniques for Association Rule Mining

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Abstract: Fast and complete retrieval of individual customer needs and "to the point" product offers are crucial aspects of customer satisfaction in todays' highly competitive banking sector. Growing number of transactions and customers have excessively boosted the need for time and memory in market basket analysis. In this paper, sampling process is included into analysis aiming to increase the performance of a product offer system. The core logic of a sample, is to dig for smaller representative of the universe, that is to generate accurate association rules. A smaller sample of the universe reduces the elapsed time and the memory consumption devoted to market basket analysis. Based on this content; the sampling methods, the sampling size estimation techniques and the representativeness tests are examined. The technique, which gives complete set of association rules in a reduced amount of time, is suggested for sampling retail banking data.

1 INTRODUCTION

Today's highly competitive sales and marketing conditions force companies to have a better understanding of their customers' needs. The strategic sales and marketing decisions, which are based on the customers' purchasing profile, succeed and increase the profitability of the companies.

Market basket analysis is a fundamental data mining technique to identify the customers' behaviors. It is used to reveal the hidden patterns in the customers' transactions and to mine the associations among products, that are often bought together. Conventional market basket analysis consists of clustering and association mining. Clustering is executed in order to create groups of customers with similar marketing behaviors. Association mining takes place to figure out patterns behind these behaviors and to understand the current purchasing behavior of look alike customers.

However, analysis on large-scale databases becomes unaffordable due to time and memory consumptions. In order to improve the memory consumption, early studies introduced new association mining algorithms (Hidber, 1999; Pei et al., 2000; Hipp et al., 2000; Zhang et al., 2008; Pei et al., 2007; Zaki and Hsiao, 2002). Other studies focused on sampling for association mining so as to reduce both time and memory consumptions (Zaki et al., 1997; Chakaravarthy et al., 2009; Toivonen et al., 1996; Riondato and Upfal, 2012).

In this paper, sampling for association mining is included into the market basket analysis. Based on the assumption of "a good representative subset of the transactions would not have any association rule loss", mining process is executed on the sample rather than the universe.

To materialize this objective, this study first investigates different sampling methods and their parameters. We have identified sample size as the most important parameter, which impacts the representativeness of a sample and the time consumption of entire mining process. To measure the success of representativeness of a sample, two statistical tests have been used. Along with the results of the tests, we have implemented two sampling methods by covering several different techniques to specify the optimal sample size. In addition, we tracked loss of association rules in samples.

This paper focuses particularly on the sampling methods and implementations for association mining process in Section 2. The section of 2.1 defines the sampling methods. Section 2.2 gives a formal definition of sample size estimation techniques for association rule mining. Section 3 covers the implementation of statistical tests for sample representativeness. Dataset and test scenario are explained in Section 4. Samples are compared in terms of the representativeness tests and derived association rules. A discussion

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of results and conclusions can be found in Section 5.

2 SAMPLING

Sampling is a statistical data selection method that creates a representative subset of the universe. It is mainly utilized whenever access to universe transactions is impossible or computations on the universe are resource-intensive. Association mining is a time and memory consuming process, which makes the sampling essential for efficient use of memory and time.

Early studies about sampling for association mining show that there exists a sample which is similar to the universe (Mannila et al., 1994). The subsequent studies concentrate on finding a lower bound for the sample size from different perspectives (Zaki et al., 1997; Chakaravarthy et al., 2009; Toivonen et al., 1996; Riondato and Upfal, 2012).

The main concern of association mining is the representativeness of a sample. Loss of any association rule is undesirable for a successful sample, hence the patterns existing in the universe should remain in the sample as well. Therefore, the sampling methods and the sampling size estimation techniques play a significant role. Section 2.1 and Section 2.2 detail the discussion about sampling methods and sampling size estimation techniques, respectively.

2.1 Sampling Methods

Sampling methods, which is the first major factor during sampling, specify the procedure to be followed during the transaction selection. The methods are divided into 2 groups, namely *probability sampling* and *non-probability sampling*.

Probability sampling methods involve random selection of transactions, whereas non-probability methods do not. Randomness ensures that all transactions in the universe have a chance of selection. Non-probability sampling methods leave some transactions out of coverage. Being not able to cover all transactions in the universe, non-probability sampling methods are not eligible for association mining. Therefore, in this study, only the probability sampling methods are considered.

Assume *U* is the universe where |U| = N. A sample *s* with |s| = n can be generated by utilization of the following probability sampling methods;

- *Simple random sampling:* All transactions have same the probability of being chosen. *n* many transactions are selected from the universe randomly.

- *Stratified random sampling:* Having a categorization in the universe, this method could be preferred. Assume there exist *m* categories with *N_i* transactions, then *n_i* transactions are randomly selected from *i*th category, where

$$n_i = \frac{N_i * n}{N} \tag{1}$$

for all $i \in [0, m-1]$. All transactions in a given category have the same probability of being chosen.

- *Systematic sampling:* All transactions have the same probability of being chosen. Transactions in the universe are sorted and divided into *k* groups, where

$$k = \frac{N}{n},\tag{2}$$

the ratio between universe size and sample size. A random number $i \in [0, n-1]$ is generated as an index for each group. Every i^{th} transaction from k groups is selected for placement into sample.

Cluster sampling: Homogeneous clusters of universe are formed, each of which has *n* transactions. A random number *j* ∈ [0, *k* − 1] is generated, where

$$k = \frac{N}{n},\tag{3}$$

the ratio between universe size and sample size. The j^{th} cluster is designated as sample.

- *Multistage sampling:* In the first stage, homogeneous clusters of universe are formed. Later, a subset of the clusters is selected by simple random sampling. In the last stage, *n* transactions are selected from the subset of clusters using simple random sampling again. All transactions in a given cluster have the same probability of being chosen.

The applicability of the methods depends on the properties of the universe. The dataset for association rule mining consists of binary values. Each value shows the existence of relation between the customers and the products. The customers and the products are represented by rows and columns, respectively.

Simple random sampling is an exceptional and easy to apply method, which is independent of the dataset. Similarly, stratified random sampling can be executed for association rule mining in large databases. It is useful when the representation of the subgroups is crucial. The strata are generated by combinations of products. For instance, four strata are created from two products, say *A* and *B*. These strata are only *A* holders, only *B* holders, both *A* and *B* holders and neither *A* nor *B* holders. However, sampling methods such as systematic sampling require a numerical value for sorting. The determination of this value is a crucial task for a correct sampling. In our case, this method cannot be applied due to lack of numerical values in the universe dataset.

Cluster sampling and multistage sampling use "natural" clusters such as geographical areas. These clusters must be homogeneous. In case where no natural homogeneous clusters found such as customerproduct ownership data including only binary values, these methods cannot be applied either.

2.2 Sampling Size Estimation Techniques

The second major factor for sampling is the estimation of the sample size. There are several sample size estimation techniques in the literature. However, in this study, we mainly focused on the techniques specialized on association mining (Zaki et al., 1997; Chakaravarthy et al., 2009; Toivonen et al., 1996; Riondato and Upfal, 2012).

Association mining can be realized in two steps, namely frequent itemset (FI) discovery and association rule (AR) generation (Agrawal et al., 1993). A randomly selected sample contains statistical errors, particularly in support and confidence calculations. Error is calculated at either FI discovery step or AR generation step. Moreover, they are compared by the corresponding value of the universe either absolutely or relatively. The classification of the sampling size estimation techniques depends on the type of the error and the step where the error has emerged (Riondato and Upfal, 2012).

Table 1: Sampling size estimation techniques.

Technique	Туре	Formula
Zaki	FI _{abs}	$\frac{-2\ln(1-\gamma)}{\Theta\delta^2}$
Toivonen	FI _{abs}	$\frac{1}{2\epsilon^2}\ln\frac{2}{\delta}$
Chakaravarthy	FI _{abs}	$rac{24}{(1-\epsilon)\epsilon^2\Theta} (\Delta + 5 + \ln rac{4}{(1-\epsilon)\Theta\delta})$
Chakaravarthy	AR _{abs}	$rac{48}{(1-\epsilon)\epsilon^2\Theta}(\Delta+5+lnrac{5}{(1-\epsilon)\Theta\delta})$
Riondato	FI _{abs}	$\frac{4c}{\epsilon^2}(v+\ln\frac{1}{\delta})$
Riondato	FI _{rel}	$\frac{4(2+\varepsilon)c}{\varepsilon^2(2-\varepsilon)\Theta} \left(\nu \ln \frac{2+\varepsilon}{\Theta(2-\varepsilon)} + \ln \frac{1}{\delta}\right)$
Riondato	AR _{abs}	$\frac{c}{\eta^2 p} \left(\nu \ln \frac{1}{p} + \ln \frac{1}{\delta} \right)$
Riondato	AR _{rel}	$\frac{c}{n^2 p} \left(v \ln \frac{1}{p} + \ln \frac{1}{\delta} \right)$

For the sake of brevity, let us give parameters used in the techniques;

ε: upper bound of absolute/relative error,

 δ : failure probability in FI discovery/AR generation step,

- Θ : minimum support of FI,
- γ : minimum confidence of AR,
- Δ : maximum transaction length,
- η: function of Θ , γ and ε,
- *p*: function of η and Θ ,
- v: d-index of the universe,
- c: constant.

Detailed explanation about these parameters and proofs of approximations can be found in the studies (Zaki et al., 1997; Chakaravarthy et al., 2009; Toivonen et al., 1996; Riondato and Upfal, 2012; Löffler and Phillips, 2009; Har-Peled and Sharir, 2011; Mannila et al., 1994). Studied sample size estimation techniques according to their classification are as follows;

2.2.1 FI - Absolute Error

The techniques in this group aim to control the error in support values occurred at FI discovery step. The absolute error is

$$error_{abs} = |supp_s - supp_u|, \tag{4}$$

where $supp_s$ and $supp_u$ denote the support of a given itemset calculated from the sample and the universe, respectively. These techniques are labeled as FI_{abs} in Table 1.

Zaki, Toivonen, Chakaravarthy and Riondato techniques utilize Chernoff bounds theorem for sample size estimation. Each technique considers a different random variable that is generated from the dataset. Zaki concentrates on the expected support of frequent itemsets, whereas Toivonen deals with the absolute support error of the frequent itemsets.

Both Chakaravarthy and Riondato study ε -close approximation of the frequent itemsets using absolute support errors. Chakaravarthy considers the longest transaction length in the universe, whereas Riondato exploits the d-index of the universe. The d-index relies on the VC dimension of the dataset, which gives information about the transactions in the universe (Vapnik and Chervonenkis, 1971). Detailed proofs can be found in the studies of (Zaki et al., 1997; Chakaravarthy et al., 2009; Toivonen et al., 1996; Riondato and Upfal, 2012).

2.2.2 FI - Relative Error

The technique in this group aims to control the error in support values occurred at FI discovery step. The relative error is

$$error_{rel} = \frac{|supp_s - supp_u|}{|supp_u|} \tag{5}$$

where $supp_s$ and $supp_u$ denote the support of a given itemset calculated from the sample and the universe, respectively. These techniques are labeled as FI_{rel} in Table 1.

Riondato technique uses Chernoff bounds theorem for sample size estimation. The d-index of the universe is exploited for a relative ε -close approximation of the frequent itemsets. The relative errors occurred in the approximations are considered in the computations. Detailed proof can be found in the study of (Riondato and Upfal, 2012).

2.2.3 AR - Absolute Error

The technique in this group aim to control the error in confidence values occurred at AR generation step. The absolute error is

$$error_{abs} = |conf_s - conf_u|$$
 (6)

where $conf_s$ and $conf_u$ denote the confidence of a given rule calculated from the sample and the universe, respectively. These techniques are labeled as AR_{abs} in Table 1.

Chernoff bounds theorem on absolute confidence errors of an ε -close approximation of the association rules is executed by both of the techniques. Chakaravarthy considers the longest transaction length in the universe, whereas Riondato exploits d-index of the universe. η and *p* parameters are functions of Θ , γ and absolute ε . Detailed proofs, the definitions of η and *p* parameters can be found in the studies of (Chakaravarthy et al., 2009; Riondato and Upfal, 2012).

2.2.4 AR - Relative Error

The technique in this group aims to control the error in confidence values occurred at AR generation step. The relative error is

$$error_{rel} = \frac{|conf_s - conf_u|}{|conf_u|} \tag{7}$$

where $conf_s$ and $conf_u$ denote the confidence of a given rule calculated from the sample and the universe, respectively. These techniques are labeled as AR_{rel} in Table 1.

Riondato technique utilizes Chernoff bounds theorem. Relative ε -close approximation of association rules by exploiting d-index of the universe is studied. The d-index relies on the VC dimension of the dataset, which gives information about the transactions in the universe (Vapnik and Chervonenkis, 1971). η and pparameters are functions of Θ , γ and relative ε . The relative errors occurred in the approximations are considered in the computations. Detailed proof and the definitions of η and p parameters can be found in the study of (Riondato and Upfal, 2012).

3 REPRESENTATIVENESS TESTS

Application of the sampling method and the sampling size estimation technique result into a statistical sample of the universe. The assumption under utilization of sampling in association mining is that, the sample is similar to universe in terms of the associations (Zaki et al., 1997; Chakaravarthy et al., 2009; Toivonen et al., 1996; Riondato and Upfal, 2012). Thus, the sample can be mined for associations instead of the universe.

Statistical hypothesis tests are used in order to determine the probability that a given hypothesis is true. In this paper, hypothesis tests are realized in order to prove the dissimilarity between the sample and the universe. These tests contain two hypotheses, which are opposite to each other, namely *null hypothesis* and *alternative hypothesis*. The null hypothesis is

$$H_0: s_i = U_i \forall i, \tag{8}$$

which assumes the sample s and the universe U have same frequencies for all categories. The alternative hypothesis is

$$H_1: s_i \neq U_i \exists i, \tag{9}$$

which assumes the sample *s* and the universe *U* differ in some frequencies for some categories. The assessment of the truth of H_0 is realized with tests, such as χ^2 and Kolmogorov-Smirnov (K-S). Each test outputs a test statistics. These statistics are converted into *p* values in order to compare with the acceptable significance level α . If $p < \alpha$, then H_0 is rejected at the given level of significance. Otherwise, test has no result, i.e. there is no sufficient evidence to prove/disprove the similarity between the sample and the universe. For this work, the significance level is assumed to be $\alpha = 0.05$. The computed *p* values are compared with 0.05, in order to reject the null hypothesis.

Statistics are computed from a different point of view in both of the tests. χ^2 test computes the average deviance in the frequencies, whereas K-S test computes the maximum deviance in the cumulative frequencies. χ^2 test is sensitive to sample size and small frequencies (Agresti, 1996). In other words, a small sample with small frequencies result into bias in test results. In K-S test, type II error rates are greater than χ^2 test (Durbin, 1973). Thus, there is a tendency towards the rejection of the null hypothesis. Therefore, the *p* values calculated from each test, may result into different values. Due to these drawbacks, both of the tests are conducted for comparison of the results.

4 EXPERIMENTATIONS

Tests are conducted on real customer-product ownership data. Original bank dataset contains 143 products, that are classified into 10 product groups according to the hierarchy defined by the bank. Aiming to speed up the test phase, product groups are mined for associations instead of all products. Universe dataset contains 1,048,575 customers and product group ownership statuses. Rows and columns represent customers and product groups, respectively.

Sample size estimation technique parameters are taken as accuracy $\varepsilon = 0.04$, failure probability $\delta = 0.07$, minimum support $\Theta = 0.02$ and minimum confidence $\gamma = \{0.06, 0.1, 0.14\}$. Accuracy and failure probability are chosen according to the study of Riondato, whereas minimum support is chosen based on the universe dataset (Riondato and Upfal, 2012). Minimum confidence varies as in the study of Riondato for fixed accuracy, failure probability and minimum support (Riondato and Upfal, 2012). Same minimum support and minimum confidence values are used for association mining process.

Simple random sampling and stratified random sampling methods were adopted. Simple random sampling draws each transaction with equal probabilities. In stratified random sampling, 4 categories are created. Two most common product groups among customers are discovered, say *A* and *B*. Combination of these two product groups are built, namely only *A* owners, only *B* owners, both *A* and *B* owners and none of *A* and *B* owners. The ratios of these categories are preserved in the samples as in the universe.

R programming language is used for code development. The following steps are followed during test phase;

- 1. Estimation of sample sizes,
- 2. If the technique estimates a smaller size than the universe size, then 10 different samples are created using simple random sampling and stratified random sampling methods for each technique,
- 3. Analysis of χ^2 and K-S tests for representativeness of the samples,
- 4. Discovery of FIs and generation of ARs from universe and sample by using *arules* package of R programming language,
- Comparison of results and calculation of error values,
- 6. Comparison of elapsed time during AR generation from universe with total time spent on sample creation and AR generation from sample.

4.1 Sample Size

Using techniques presented in Section 2.2, we have calculated sample sizes. Table 2 shows sample size estimations for varying γ values which is the main parameter impacting the sample size. Sample sizes estimated by Toivonen, Chakaravarthy FI_{abs}, Chakaravarthy AR_{abs}, Riondato FI_{abs} ve Riondato FI_{rel} do not differ with γ values, since γ is not taken as a parameter in these formulae.

Table 2: Sample size estimations varying with minimum confidence.

Technique	Туре	$\gamma =$	$\gamma =$	$\gamma =$
		0.06	0.10	0.14
Zaki	FIabs	3867	6585	9426
Toivonen	FIabs	1047	1047	1047
Chakaravarthy	FIabs	$\approx \! 15M$	$\approx \! 15M$	$\approx \! 15M$
Riondato	FIabs	9574	9574	9574
Riondato	FI _{rel}	$\approx \! 15M$	$\approx \! 15M$	$\approx \! 15M$
Chakaravarthy	AR _{abs}	$\approx 30 \text{M}$	$\approx 30 \text{M}$	$\approx 30 \text{M}$
Riondato	AR _{abs}	15057	47005	96859
Riondato	AR _{rel}	$\approx 5M$	$\approx 5M$	$\approx 5M$

Chakaravarthy FI_{abs} , Chakaravarthy AR_{abs} , Riondato FI_{rel} and Riondato AR_{rel} techniques offer bigger sizes than universe size (1048575). As one of our main concern is to find the optimal sample size, a sample with a greater size than the actual universe is an undesired outcome. Hence the techniques stated above have been left out of further investigations. Table 2 presents the results of sample size estimations.

4.2 *p* **Value**

For each sample size presented in Table 2 that are smaller than the universe size, we have created samples using simple random sampling and stratified random sampling methods. We have drawn 10 different samples in order to minimize error arising from randomness of sampling methods.

We adopted p value for testing null hypothesis. We determined $\alpha = 0.05$ as significance level of the tests. If p values are less than the significance level, null hypothesis, i.e. the sample is similar to the universe, is rejected.

We present average p values calculated from χ^2 and K-S tests in Table 3 for the samples obtained from simple random sampling and Table 4 summarizes the same test results for the samples obtained from stratified random sampling.

The major interpretation of p values to understand the representativeness of a sample is that, the p values are greater than the significance level. Thus, there is

Technique	Туре	$\gamma =$	$\gamma = 0.06$		$\gamma = 0.10$		$\gamma = 0.14$	
		χ^2	K-S	χ^2	K-S	χ^2	K-S	
Zaki	FI _{abs}	0.409	0.364	0.605	0.636	0.496	0.313	
Toivonen	FI _{abs}	0.429	0.365	0.643	0.408	0.589	0.339	
Riondato	FI _{abs}	0.437	0.583	0.531	0.670	0.458	0.493	
Riondato	AR _{abs}	0.425	0.328	0.558	0.499	0.612	0.575	

Table 3: Average p values for simple random samples computed from χ^2 and K-S tests.

Table 4: Average p values for stratified random samples computed from χ^2 and K-S tests.

Technique	Туре	$\gamma =$	$\gamma = 0.06$		$\gamma = 0.10$		$\gamma = 0.14$	
		χ^2	K-S	χ^2	K-S	χ^2	K-S	
Zaki	FI _{abs}	0.606	0.417	0.697	0.525	0.576	0.456	
Toivonen	FI _{abs}	0.619	0.475	0.501	0.525	0.477	0.264	
Riondato	FI _{abs}	0.624	0.504	0.710	0.428	0.505	0.640	
Riondato	AR _{abs}	0.618	0.451	0.692	0.649	0.577	0.481	

no sufficient evidence to disprove statistical similarity between the universe and the samples.

p values obtained by χ^2 and K-S tests are not equivalent because of the reasons explained in Section 3. Besides, we could not observe any proportionality between sample sizes and *p* values. For the samples with constant sizes but with varying γ values, the application of the Toivonen technique has provided unstable *p* values.

We have tested both simple random sampling and stratified random sampling methods under different techniques but we cannot observe any significant difference between results.

4.3 Absolute Support and Confidence Errors

Discovery of FIs and generation of ARs are realized by Apriori algorithm. The absolute errors in support and confidence values are measured for FIs and ARs respectively. Zaki and Toivonen techniques miss some FIs and ARs that exist in the universe. Corresponding p values of these techniques did not give sufficient information to disprove the similarity. However, using absolute errors, we noticed that these two sample size estimation techniques do not give satisfactory results. The ineffectiveness of these methods depends on their small sample sizes.

Since loss in association rules is undesirable, these two techniques are interpreted as ineligible for sampling and they are not subject to further testing. Missing support and confidence values are taken as 0. Average support and confidence errors for simple random samples (respectively for stratified random samples) are shown in Figure 1 (respectively in Figure 2) for varying minimum confidence. As shown in the figures, Riondato's techniques present better results than other techniques in terms of support and confidence errors.

In compliance with our expectations, the confidence errors are high whenever support error is high. In addition, there is a relation between the sample size and the errors. The bigger the sample sizes, the smaller the errors (See Table 2 for sample size variation).

Comparing the results of simple random sampling and stratified random sampling methods, we did not observe any remarkable difference.



Figure 1: Average support and confidence errors for simple random samples for varying minimum confidence γ .



Figure 2: Average support and confidence errors for stratified random samples for varying minimum confidence γ .

4.4 Time Consumption

We present time consumptions for both universe and sample until the end of AR generation in Figure 3 and Figure 4. Tests are conducted on a PC with 3.2 G Hz i5 processors and 8 GB RAM.

For universe, only AR generation time is given, whereas for sampling techniques, average total time for sample size estimation, simple random sampling or stratified random sampling and AR generation is given. As expected, for all γ values, sampling techniques performed at least approximately six times better than universe. We anticipate that time performance will be more visible when 143 products are included in the sampling process rather than just 10 groups of product.



Figure 3: Time consumptions (in seconds) until AR generation for simple random samples.



Figure 4: Time consumptions (in seconds) until AR generation for stratified random samples.

5 CONCLUSIONS AND FUTURE WORKS

We targeted creating a representative small size sample out of universe and mining it for association rule retrieval. The initiative for sampling is to minimize memory and time consumptions. For this purpose, sampling size estimation techniques specialized on association mining, sampling methods and representativeness tests are investigated and applied in this study. Samples are created by utilizing simple random sampling and stratified random sampling methods to crosscheck the results. For each technique satisfying size condition, 10 samples are created. The rationale behind multiple samples for a given technique is to minimize noise introduced by random sampling methods. Techniques are tested with varying minimum confidence values. The choice of parameter values is based on the study of Riondato. The very first interpretation about techniques is that not all of them are applicable for any size of universe. We observed 3 different indicators in order to compare the results.

First, we examined p value as an a-priori indicator of representativeness. According to p values, we could not reject any sample because of representativeness. However, when we compare FIs and ARs from the universe and the sample, we realized that some techniques miss some FIs and ARs, leading to information loss. For instance, sample sizes calculated by Toivonen technique are not rejected because of representativeness according to p values for $\gamma = 0.06$. However, it misses some FIs and ARs. Hence, we conclude that traditional statistical methods are not suitable tools for testing representativeness of samples created for association mining. One of our contributions is the identification of the unavailability of the statistical tests to check the representativeness of sampling for association mining.

In addition, observations in absolute errors in confidence and support helped us to identify sample size estimation techniques leading to loss of FIs and ARs. We noticed that use of these techniques for association mining is not convenient.

Examining the absolute errors, we verified that the bigger the sample size, the smaller the absolute errors. Riondato FI_{abs} and AR_{abs} return the smallest absolute errors due to their high sample size. We could not find any threshold of acceptance for errors identified in the studies of this domain. In future works, we aim to determine this threshold.

Besides, time elapsed during AR generation from the universe is compared with total time elapsed during sample size estimation, creating samples with simple random sampling or stratified random sampling method and AR generation from the sample. Each of the techniques performed better than universe in terms of time consumption. According to the absolute errors, Riondato FI_{abs} and Riondato AR_{abs} techniques are the best performers. When smaller sample size and less time consumption criteria are taken in concern, Riondato FI_{abs} is the leading sample size estimation technique. Among several different sample size estimation techniques, we identified Riondato FI_{abs} to be the most suitable technique for our retail banking data.

Dataset contains retail bank customers and their product group ownership information. Rather than the individual products owned by the customers, the groups of these products are taken into consideration. The main reason for this decision is to eliminate sparsity on the dataset and speed up the test phase. Use of product groups has led to small time consumptions even on the universe. Even though duration gain seems to be in the order of seconds, bigger gains can be obtained if larger product sets are tested. For next studies, dataset will be expanded and robustness check will be done with an alternative dataset.

Moreover, systematic, cluster and multistage sampling methods can be applied during construction of association rule mining data. For instance, the customers which will be subjects of ARM can be drawn according to their clusters (e.g. geographical areas). Our research focuses on the association rule mining data (including binary values). Extraction of this data from customers' dataset will be examined in further studies.

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