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STUDY ON IMPLEMENTING ASSOCIATION RULE MINING IN PARTICLE SWARM OPTIMIZATION

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ABSTRACT

There is a great need to discover association rules which are an important database mining problem. In the area of association rule mining is focused on improving computational efficiency in previous research. Determination of the threshold value of support and confidence affect the quality of association rule mining up to great extent. This paper gives attention to effectiveness of algorithm for association rule mining in order to improve computational efficiency as well as to automatically determine suitable threshold values. The Particle Swarm Optimization algorithm first search for the optimum fitness value of each particle and finds corresponding support and confidence as minimal threshold values after the data are transformed into binary values. In this paper the particle swarm optimization algorithm compared with a genetic algorithm.

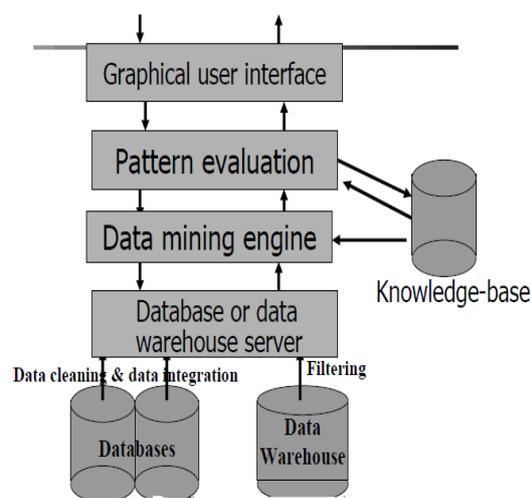
KEYWORDS

Association Rule Mining, Data Mining, Particle Swarm Optimization, Genetic Algorithm.

I. INTRODUCTION

Data Mining is process which finds useful patterns from large amount of data analysis is that the data mining is mine information and discover knowledge on the premise of no clear assumption.[1]. Generally data mining process is composed by preparation of data, data cleaning, data reduction and transformation, data mining, pattern evaluation and Knowledge presentation. Fig[1] shown the process of data mining.

FIGURE 1: PROCESS OF DATA MINING



Association rule mining is one of model among data mining several models. Association rule mining is used to searches for relationship between variables in database. Apriori algorithm is one of famous and oldest algorithm for discover the association rules. It consists of many modified algorithms to improving computational efficiency. The Support and Confidence is effect the quality of result and also it's used to decision making purpose. But thus previous algorithm doesn't consider that main two parameters. So this study mainly focused to improving algorithm can find minimum threshold value for support and confidence.

II. LITERATURE REVIEW

2.1 ASSOCIATION RULE MINING:

Association Rule analysis is useful for discovering interesting relationship hidden in large data sets. The uncovered relationship can be represented in the form of association rules. Agarwal et al was introducing association rules in 1993. Retailers are interested in analyzing the data to learn about the purchasing behavior of their customers. Such valuable information can be used to support a variety of business-related application such as marketing promotions, inventory management, and customer relationship management. An association rule is an implication expression of the form $X \rightarrow Y$, where X and Y are disjoint item sets. The strength of an association rule can be measured in terms of its Support and Confidence. Support determines how often a rule is applicable to a given data set, while confidence determines how frequently items in Y appear in transaction that contains X.

$$\text{Support, } s(X \rightarrow Y) = \frac{\delta(XUY)}{N}$$

$$\text{Confidence, } c(X \rightarrow Y) = \frac{\delta(XUY)}{N}$$

2.2 ASSOCIATION RULE ALGORITHM

2.2.1. Apriori Algorithms: The classical apriori algorithm as suggested by Agrawal et al in [1993] is one of the most important data mining algorithms. It is an influential algorithm for mining frequent item sets for Boolean association rules. Frequent Item sets: The sets of item which has minimum support. Apriori property: Any subset of frequent item set must be frequent. Joint operation: To find L_k , a set of candidate K-Item sets is generated by joining L_{k-1} with itself.

2.2.2. FP-Growth Algorithm: J.Han et al [2000], it's for complete set of frequent patterns, by pattern fragment growth. Efficiency of mining is achieved with three techniques: 1. A large database is compressed into a condensed smaller data structure, FP-tree which avoids the costly, repeated database scans, 2. FP-tree based mining adopts a pattern – fragment growth method to avoid costly generation of large number of candidate sets, 3. A partitioning based, divide-and-conquer method is used to decompose the mining task into a set of smaller tasks for mining confined patterns in conditional database, which dramatically reduces the search space.[2]

```

Join Step:  $C_k$  is generated by joining  $L_{k-1}$  with itself

Prune Step: Any  $(k-1)$ -itemset that is not frequent cannot be a subset of a frequent  $k$ -itemset

 $C_k$ : Candidate itemset of size  $k$ 
 $L_k$ : frequent itemset of size  $k$ 
 $L_1 = \{\text{frequent items}\};$ 
for  $(k=1; L_k \neq \emptyset; k++)$  do begin
 $C_{k+1}$  = candidates generated from  $L_k$ ;
for each transaction  $t$  in database do
increment the count of all candidates in  $C_{k+1}$  that are contained in  $t$ 
 $L_{k+1}$  = candidates in  $C_{k+1}$  with min. support
end
return  $\cup_k L_k$ ;

```

2.2.3. Dynamic FP-Growth Algorithm: C.Gyorodi et al [2003], It has focused in improving the FP-Tree algorithm construction based on two observed problems: 1. The resulting FP-Tree is not unique for the same "Logic" database. 2. The process needs two complete scans of the database. The above problems solved by G.Gyorodi et al. An important feature in this approach is that it's not necessary to rebuild the FP-Tree when the actual database is updated. It's only needed to execute the algorithm again taking into consideration the new transaction and stored FP-Tree.[3]

2.2.4. Partition Algorithm: A.Savasere et al [1995], Partition algorithm based on apriori algorithm, but it requires only two complete scans over the database. The Partition algorithm divided into two phases: 1. the database is divided into a number of non-overlapping partitions and frequent item sets local to partition are generated for each partition. The database is scanned completely for the first time. 2. Local frequent item sets from each partition are combined to generate global candidate item sets. Then the database is scanned second item to generate global frequent item sets [4].

2.2.5. DIC Algorithm: S.Brin et al [1997], DIS is a further variation of the Apriori-algorithm. DIC is a softens the strict separation between counting and generating candidates. Whenever a candidate reaches minsup that is even when this candidate has not yet "seen" all transactions, DIC starts generating additional candidates based on it. For that purpose a prefix-tree is employed. In contrast to the hash tree, each node-leaf node or inner node of the prefix-tree is assigned to exactly one candidate respectively frequent item set. In contrast to the usage of a hash tree that means whenever it reach a node it can be sure that the item set associated with this node is contained in the transaction furthermore interlocking support determination and candidate generation decrease the number of database scans.[5]

2.2.6. Pincer Search Algorithm: This algorithm was proposed by Dao. I et al [1997].The algorithm was uses both the top-down and bottom –up approaches to Association rule mining. It is a slight modification to original Apriori algorithm. The main search direction is bottom -up expect that it conducts simultaneously a restricted top-down search. [6]

2.2.7. Pincer Search Algorithm: The concept of PSO was first suggested by Kennedy and Eberhart in 1995[5]. Particle swarm optimization (PSO) is inspired by the social behavior observed in flocks of birds and schools of fish. In nature, there is a leader who leads the bird or fish group to move, as illustrated in Fig. 2. Most members of the group follow the leader. In PSO, a potential solution to the considered problem is represented by a particle, similar to the individuals in the bird and fish group. Each particle travels in the solution space and attempts to move toward a better solution by changing its direction and speed based on its own past experience and the information from the current best particle of the swarm. [7]

2.2.8. The procedure of PSO is described as follows:

A. Particle initialization:

An initial swarm of particles is generated in search space. Usually, the population size is decided by the dimension of problems.

B. Velocity and position update:

In each iteration, a new velocity value for each particle is calculated based on its current velocity, the distance from its previous best position, and the distance from the global best position. The new velocity value is then used to calculate the next position of the particle in the search space. The particle's velocity and position are dynamically updated as follows:

$$V_{id}^{new} = w \times V_{id}^{old} + c_1 \times rand \times (P_{id} - X_{id}^{old}) + c_2 \times rand \times (P_{gd} - X_{id}^{old}),$$

$$X_{id}^{new} = X_{id}^{old} + V_{id}^{new}.$$

The new velocity of a particle, V_{newid} , is updated by taking into consideration of the particle's previous velocity, V_{oldid} , and previous position, x_{oldid} . $w = [0.5 + rand/2]$ is an inertia weight and $rand$ is a uniformly generated random number between 0 and 1. The cognition parameter, c_1 , and social parameter, c_2 , are acceleration coefficients that are conventionally set to a fixed value 0–2. P_{id} is the previous individual best position of this particle and P_{gd} is the current global best position then calculates the new position of the particle, x_{newid} . [8]

- a) Both particle's position and the global best position are far from the optimum and the particle velocity is low compared to its distance to the optimum [9].
- b) Global best position is close to the optimum and the particle position is far from them resulting in a small improvement region and a large next position region.

C. Evaluation and update of best locations:

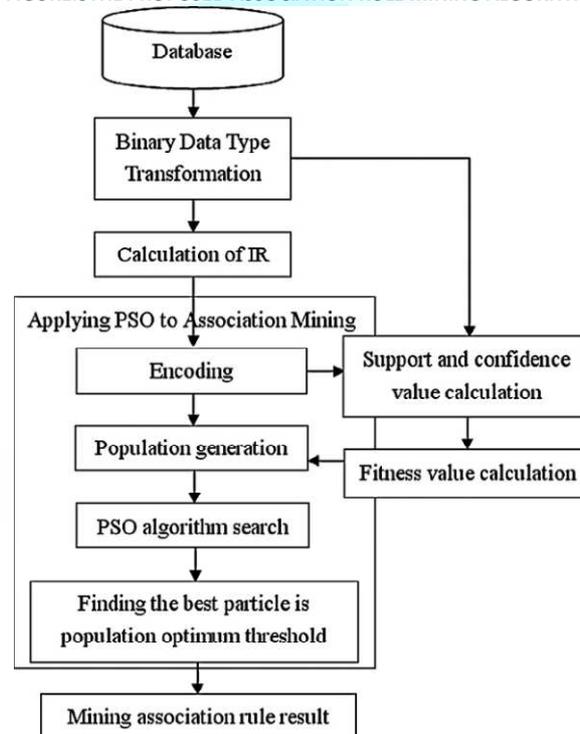
The fitness value of each particle is calculated by the objective function. The values of P_{id} and P_{gd} are then evaluated and replaced if better particle best position or global best position is obtained.

D. Termination:

Steps (2) and (3) are repeated iteratively until the termination condition is met.

III METHODOLOGY

FIGURE:3 THE PROPOSED ASSOCIATION RULE MINING ALGORITHM



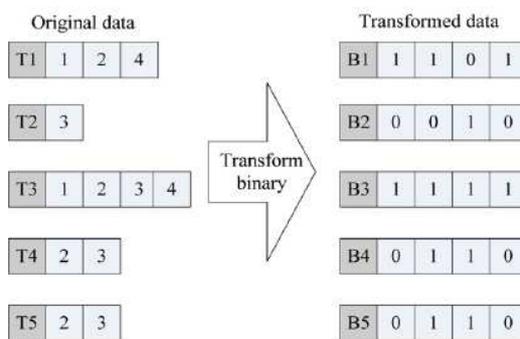
3.1. The proposed algorithm:

The proposed algorithm comprises two parts, preprocessing and mining. The first part provides procedures related to calculating the fitness values of the particle swarm. Thus, the data are transformed and stored in a binary format. In the second part of the algorithm, which is the main contribution of this study, the PSO algorithm is employed to mine the association rules. First, it proceeds with particle swarm encoding, this step is similar to chromosome encoding of genetic algorithms. The next step is to generate a population of particle swarms according to the calculated fitness value. Finally, the PSO searching procedure proceeds until the stop condition is reached, which means the best Particle is found. The support and confidence of the best particle can represent the minimal support and minimal confidence. [10] Thus, it can use this minimal support and minimal confidence for further association rule mining. Fig.3 illustrates the algorithm structure.

3.2. Preprocessing of PSO association rule mining

Binary transformation: In this transform the transaction data into binary type data, each recorded and stored as either 0 or 1 [8]. This approach can accelerate the database scanning operation, and it calculates support and confidence more easily and quickly. The transformation approach is explained by an example in Fig. 4. In Fig. 4, there are five records, say T1 to T5, in the original data. Each of these records is transformed and stored as a binary type. For instance, there are a total of only four different products in the database, so four cells exist for each transaction. Take B4 as an example, this transaction only purchased products 2 and 3, so the values of cells 2 and 3 are both "1s," whereas cells 1 and 4 are both "0s." Conversion shown in figure 6. Fig. 4: Data type transformation

FIGURE 4: DATA TYPE TRANSFORMATION



3.3. Application of PSO to association rule mining

Applying PSO to association mining is the main part of this study. We use PSO as a module to mine best fitness value. The algorithmic process is quite similar to that of genetic algorithms, but the proposed procedures include only encoding, fitness value calculation, population generation, best particle search, and termination condition. Each of the steps in the PSO algorithm and the process of generating association rules are explained as follows:

Encoding: According to the definition of association rule mining, the intersection of the association rule of item set X to item set Y (X→Y) must be empty. Items which appear in item set X do not appear in item set Y, and vice versa. Hence, both the front and back partition points must be given for the purpose of chromosome encoding. The item set before the front partition point is called "item set X," while that between the front Partition and back partition points is called "item set Y."

3.4. Fitness value calculation: The fitness value in this study is utilized to evaluate the importance of each particle. The fitness value of each particle comes from the fitness function. Here, employ the target function [11] to determine the fitness function value as shown in Eq.

$$Fitness(k) = confidence(k) \times log(support(k) \times length(k) + 1)$$

Fitness (k) is the fitness value of association rule type k. Confidence (k) is the confidence of association rule type k. Support (k) is the actual support of association rule type k. Length (k) is the length of association rule type k. The objective of this fitness function is maximization. The larger the particle support and confidence, the greater the strength of the association, meaning that it is an important association rule.

3.5. Population generation: In order to apply the evolution process of the PSO algorithm, it is necessary to first generate the initial population. In this study, it has selected particles which have larger fitness values as the population. The particles in this population are called initial particles.

FIGURE 5: BINARY TRANSFORMATIONS

Transaction Id	Cat2	Cat3	Cat4	...	Cat34
3:483	0	0	0	0	0
3:483	0	0	0	0	0
3:661	0	0	0	0	0
5:370	0	0	0	0	0
6:570	0	0	0	0	0
10:386	0	0	0	0	0
10:424	0	0	0	0	0
14:559	0	0	0	0	0
19:553	0	0	0	0	0
20:372	0	0	0	0	0
26:658	0	0	0	0	0
28:384	0	0	0	0	0
28:654	0	0	0	0	0
30:448	0	0	0	0	0
30:495	0	1	0	0	0
30:570	0	0	0	0	0
36:616	0	0	0	0	0
36:686	0	0	0	0	0
39:485	0	0	0	0	0
39:586	0	0	0	0	0
40:466	0	0	0	0	0
41:448	0	0	0	0	0
41:636	0	0	0	0	0
43:427	0	0	0	0	0
44:470	0	0	0	0	0
44:564	0	0	0	0	0
44:585	0	0	0	0	0
44:659	0	0	0	0	0
44:720	0	0	0	0	0
45:412	0	0	0	0	0
45:524	0	0	0	0	0
49:509	0	0	0	0	0
49:567	0	0	0	0	0
49:652	0	0	0	0	0
50:561	0	0	0	0	0
50:628	0	0	0	0	0
62:552	1	0	0	0	0
64:428	0	0	0	0	0
74:417	0	0	0	0	0

Fitness value calculation is shown in figure 5

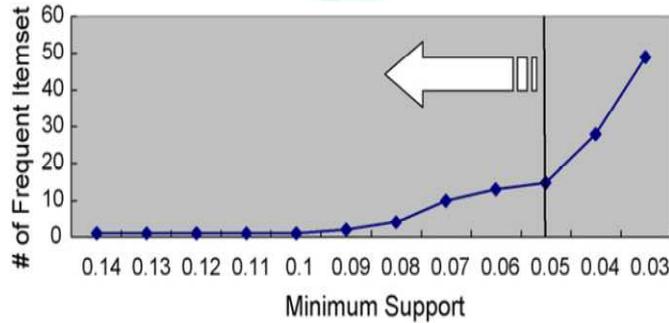
3.6. Search the best particle: First, the particle with the maximum fitness value in the population is selected as the "gbest." It designed a method to constrain this search. The constrained method is to calculate the distance between the particle's new position and all the possible particles inside the constrained range before the particle's position is updated. Definitely, the particle with the smallest distance will be selected and treated as the particle's new position. In the distance measuring function it uses traditional "Euclidean distance" as shown in Eq.

$$\text{dist}(x^n, y^m) = \sqrt{\sum_1^d (x_i^n - y_i^m)^2}$$

Where x_n is the position of the particle at nth update and y_m is the possible particle number min the constrained range. In addition, d is the dimension of the search space. The nearest possible particle is selected to be the target particle's new position. This method can prevent a particle from falling beyond the search space when its position is updated.

3.7. Termination condition: To complete particle evolution, the design of a termination condition is necessary. In this study, the evolution terminates when the fitness values of all particles are the same. In other words, the positions of all particles are fixed. Another termination condition occurs after 100 iterations and the evolution of the particle swarm is completed. Finally, after the best particle is found, its support and confidence are recommended as the value of minimal support and minimal confidence as shown in Fig[6]. These parameters are employed for association rule mining to extract valuable information. [10]

FIGURE 6: NUMBER OF HIGH-FREQUENCY ITEM SETS UNDER DIFFERENT MINIMAL SUPPORTS



IV. Model evaluation results and discussion

This section will use the database provided by Microsoft SQL Server 2000 to verify the feasibility of the proposed algorithm. A detailed discussion is provided as follows.

4.1. Experimental platform and database

This study's experiment was conducted in the environment of Microsoft Windows XP using an IBM compatible computer with Inter Pentium IV 1.60GHz and 512MB RAM. The algorithm was coded by Borland C++ Builder 6. In regard to the experimental testing database, its source was a FoodMart2000 retail transaction database embedded in a Microsoft SQL Server 2000, as illustrated in Fig. 6. Since there are different kinds of transaction databases in FoodMart2000, it only selects sales fact 1997 data table for assessment. The number of product items in this data table is 1560.

In order to effectively mine meaningful association rules, this experiment categorizes the products into groups according to the product category provided by the data table. Thus, products are classified into 34 categories, each with a corresponding product category id. In regard to data selection, 6000 customers' are randomly selected along with their corresponding transaction data at different times. After arrangement, there are a total of 12,100 transaction records for these 6000 customers.

FIGURE 7 THE DATA TABLE OF THE FOODMART2000 DATABASE

customer_id	time_id	product_category_id	product_category
5845	727	31	Snack Foods
5845	727	23	Meat
5845	727	23	Meat
5845	727	3	Beer and Wine
5845	727	29	Seafood
5846	386	14	Electrical
5846	386	15	Frozen Desserts
5846	386	19	Hygiene
5846	386	17	Fruit
5846	386	20	Jams and Jellies
5846	536	13	Eggs
5846	536	17	Fruit
5846	536	7	Canned Soup
5846	536	27	Pizza
5846	585	8	Hot Beverages
5846	585	26	Paper Products
5846	585	4	Bread
5846	585	6	Candy
5846	585	31	Snack Foods
5846	600	31	Snack Foods
5846	600	21	Kitchen Products
5846	600	2	Bathroom Products
5846	600	3	Beer and Wine
5846	600	1	Baking Goods
5846	720	15	Frozen Desserts
5846	720	18	Hardware
5847	439	34	Vegetables
5847	439	34	Vegetables

4.2. Mining results via PSO algorithm

In this experiment, every transaction record in the Food-Mart2000 has 1-13 items. After calculating the IR values, it is found that $R(1 \rightarrow 6) = 5.86822$ is the largest. Therefore, it can generate five different dimensions of encoding types for the particle swarm. They are two dimensions, 1-2, three dimensions, 1-3 and 2-3, four dimensions, 1-4, 2-4, and 3-4, and five dimensions, 1-5, 2-5, 3-5, and 4-5, and six dimensions, 1-6, 2-6, 3-6, 4-6, and 5-6. According to these five dimensions, it can implement the PSO mining process. An example of two dimensions is illustrated in Fig. 6. Since the computational results are different for each replication, a total of 30 replications are conducted in order to get the final experimental results. The results show that there are three possibilities as listed in Table 1.

Next, it can conduct the three-dimensional PSO association rule mining. The population size is 20. The computational results can be found in Appendix A. The results indicate that the maximal support value is 0.03652 for the minimal support threshold value. Because the support threshold value is smaller than 0.05, the number of high-frequency item sets mined is too large. This means that more meaningless rules are generated. In the current experiment, since the maximal support values of minimal support for three, four, five, and six dimensions are all smaller than 0.05, none of the mining results are used.

TABLE: 1 THE RESULTS OF THE TWO-DIMENSIONAL PSO ASSOCIATION RULES FOR FOODMART2000

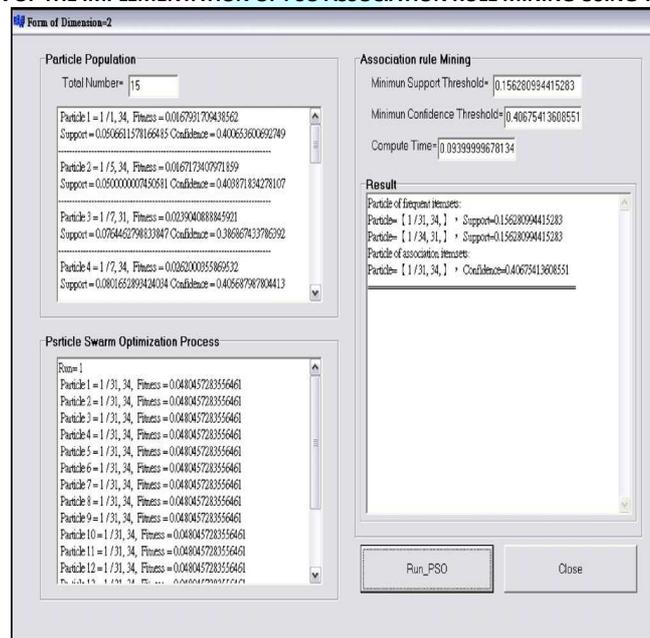
Possible results	High-frequency itemset(\geq minimal support)	Association itemset (\geq minimal confidence)
1	Minimal support = 0.15628 {Snack Foods, vegetables}	Minimal confidence = 0.40675 {Snack Foods \rightarrow Vegetables}
2	Minimal support = 0.15628 {Snack Foods, Vegetables}	Minimal confidence = 0.38497 {Snack Foods \rightarrow Vegetables} {Vegetables \rightarrow Snack Foods}
3	Minimal support = 0.09578 {Dairy, Vegetables} {Snack foods, Vegetables}	Minimal confidence = 0.40709 {Dairy \rightarrow Vegetables}
Final result	{Snack Foods, Vegetables} {Dairy, Vegetables}	Snack Foods \rightarrow Vegetables} {Vegetables \rightarrow Snack Foods} {Dairy \rightarrow Vegetables}

4.3. Performance evaluation analysis

4.3.1. Comparison of PSO and GA

Though Section 4.2 has shown that the proposed PSO algorithm can provide very promising results, further investigation is still necessary. Thus, it can compare with the genetic algorithm proposed in [9]. For both the PSO algorithm and genetic algorithm, the number of product items and the number of transaction records are 25 and 140,000, respectively. Basically, the two algorithms were implemented under the same conditions.

FIGURE: 8 A DEMONSTRATION OF THE IMPLEMENTATION OF PSO ASSOCIATION RULE MINING USING THE FOODMART2000 DATABASE



It attempt to ascertain whether or not computation time is related to the population size or number of evolutions, which are illustrated in Figs. 9 and 10, respectively. Figs. 9 and 10 clearly indicate that the proposed PSO algorithm out performs the genetic algorithm both in population size and number of evolutions. Table 2 also shows a similar outcome, showing that PSO can converge faster. This is very important as the database is very large.

4.3.2. Performance evaluation of the PSO algorithm's searching capability

This subsection intends to discuss the searching performance of the PSO algorithm. The first issue is the relationship between population size and speed. The parameter setup is as follows:

- (1) Number of dimensions: 2.
- (2) Number of data: 12,100.
- (3) Number of replications: 20.
- (4) Population size: 5, 10, 20, 30, 40 and 50 Particles.

The average running times for different population sizes are illustrated in Fig. 11. In addition, if most of the parameter setups are the same except that the size of population is 20, termination condition is 100 generations and the number of experimental replications is 50, then the relationship between the number of evolutions and running speed is presented in Fig. 12. In summary, though the proposed PSO algorithm for association mining requires more computation time with increasing population size, the increase is not significant. In addition, the number of evolutions mostly fall within the range from 1 to 10. This means that the convergence of computation is very fast. Thus, only a small number of evolutions are good enough for real application.

Furthermore, in regard to the selection of threshold value setup, this study can provide the most feasible minimal support and confidence. This dramatically decreases the time consumed by trial-and-error. Thus, the proposed PSO algorithm is better than the traditional Apriori algorithm since it does not need to subjectively set up the threshold values for minimal support and confidence. This can also save computation time and enhance performance.

FIGURE:9 RELATIONSHIP BETWEEN POPULATION SIZE AND COMPUTATION TIME FOR PSO AND GA

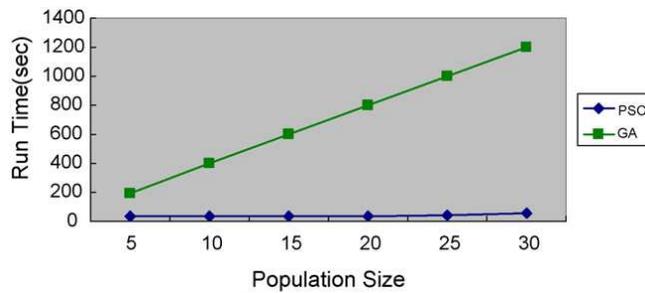


FIGURE:10 RELATIONSHIP BETWEEN NUMBER OF EVOLUTIONS AND COMPUTATION TIME FOR PSO AND GA.

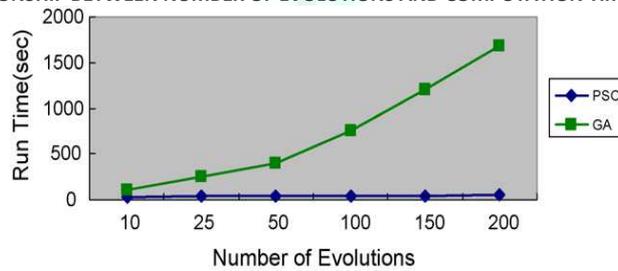


FIGURE: 11 RELATIONSHIP BETWEEN POPULATION SIZE AN COMPUTATION TIME USING THEFOODMART2000 DATABASE

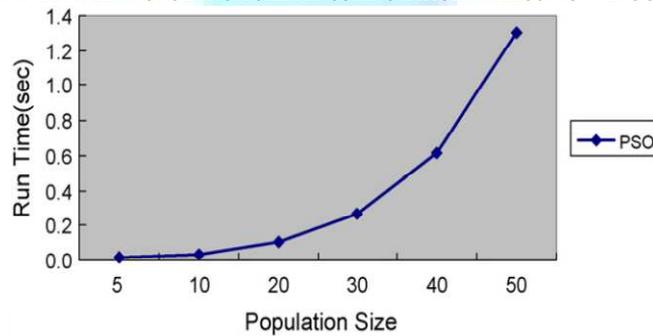
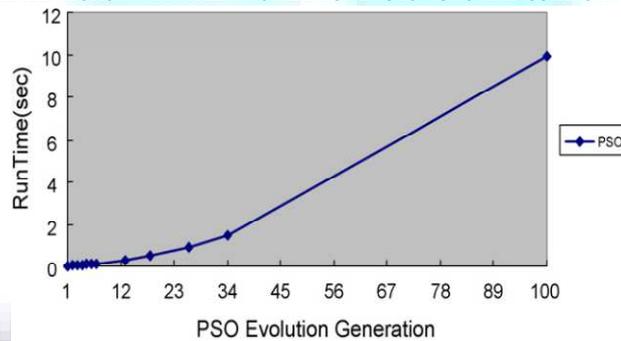


FIGURE: 12 RELATIONSHIP BETWEEN NUMBER OF EVOLUTIONS AND COMPUTATION TIME.



V. CONCLUSION

Following the development of information technology, searching for meaningful information in large databases has become a very important issue. That explains why association rules mining is the most popular technique in data mining. However, the traditional Apriori algorithm has a very critical drawback in that the minimal support and confidence is determined subjectively.

This study has demonstrated that using the PSO algorithm can determine these two parameters quickly and objectively, thus, enhancing mining performance for large databases by applying the FoodMart2000 database.

Future studies can focus on testing different updating rules. Moreover, different product items may have different importance. A weighted PSO mining algorithm could be further investigated in order to provide more practical approaches for industries.

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APPENDIX

APPENDIX A. THREE-DIMENSIONAL EXPERIMENTAL RESULTS OF FOODMART 2000

Possible result	High-frequency item set (≥minimal support)	Association item set (≥minimal confidence)
1	Minimal support = 0.03652 {Dairy, Snack Foods, Vegetables}	Minimal confidence = 0.41424 {Dairy, Snack Foods → Vegetables}
2	Minimal support = 0.03239 {Can, snack foods, vegetables} {Dairy, Snack Foods, Vegetables}	Minimal confidence = 0.4237 {Can, Snack Foods →Vegetables}
3	Minimal support = 0.02950 {Can, Snack Foods, Vegetables} {Dairy, Snack Foods, Vegetables} {Fruit, Snack Foods, Vegetables} {Jams and Jellies, Snack Foods, Vegetables} {Meat, Snack Foods, Vegetables}	Minimal confidence = 0.4204 {Can, Snack Foods →Vegetables} {Fruit, Snack Foods →Vegetables} {Jams and Jellies, Snack Foods → Vegetables}
4	Minimal support = 0.025785 {Can, Snack Foods, Vegetables} {Beverages, Snack Foods, Vegetables} {Dairy, Snack Foods, Vegetables} {Fruit, Snack Foods, Vegetables} {Jams and Jellies, Snack Foods, Vegetables} {Meat, snack Foods, Vegetables}	Minimal confidence = 0.41106 {Can, Snack Foods →Vegetables} {Beverages, Snack Foods, Vegetables} {Dairy, Snack Foods, Vegetables} {Fruit, Snack Foods →Vegetables} {Jams and Jellies, Snack Foods → Vegetables}

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