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Association Rule Generation in Data Streams using FP-Growth and APRIORI MR Algorithms

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ABSTRACT: Data stream is used for handling dynamic databases in which data can be arrived continuously, limitless and its size are very large. This situation has created a problem, i.e. to perform the mining process in these database, the existing data mining algorithms are not suitable. In order to perform mining task in data streams there is a need for development of new algorithms and techniques. By using this new algorithms and techniques we can able to perform various data mining tasks, i.e. clustering, classification, frequent pattern mining and association rule mining in data streams. Association rule mining is used to find the association between the data items which are exist in the databases. Even though, the traditional algorithms are not suitable for data streams, this paper concentrated on how to perform association rule generation task in data streams using traditional algorithms in order to find the drawbacks as well as comparing the performance of the traditional algorithms. Frequent Pattern Tree Growth algorithm and APRIORI Map/Reduce algorithms are used for generating association rules in data streams. Performance measures used in this work are execution time and number of association rules generated. From the experimental results we come to know that the performance of FP-Tree Growth algorithm is more efficient than APRIORI Map/Reduce algorithm.

KEY WORDS: Association Rules, FP-Tree Growth Algorithm, APRIORI Map/Reduce Algorithm, Rapid Miner tool, Tanagra tool.

I. INTRODUCTION

The data stream is continuous arrival of data and this data is normally dynamic in nature and its size is very huge. In this situation, it is not possible to perform data mining tasks with the traditional algorithms since those algorithms are suitable for static data bases. Hence, data stream mining needs the development of new algorithms and techniques to perform the data mining tasks. Association rules are described by finding the frequent pattern, links, relationship and the related structures among the data objects in the databases. There are two important steps in association rule mining; first step is to find the frequent data items and second step is to generate association rules from the frequent data items. This work has compared two different types of association rule mining algorithms; they are APRIORI Map/Reduce algorithm and Frequent Pattern Tree Growth algorithm. [15] The main objective of this work is to find the drawbacks of the existing algorithm when it is applied to the data streams and also finding the efficiency of the two existing algorithms APRIORI Map/Reduce and Frequent Pattern Tree Growth.

The paper is organized as follows. Section 2 provides the related works. Proposed methodology and the traditional association rule algorithms are given in Section 3. Section 4 described the experimental results. Conclusion is given in Section 5.

II. RELATED WORK

Vijayarani S et al., [17] described frequent item-set mining in data streams. Authors have used ECLAT and RARM algorithms for generating the frequent item-sets. The dataset was divided into five windows with different threshold values. The performance factors used are number of frequent items generated and execution time. From the results, the authors observed that the performance of ECLAT is more efficient than RARM.



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Vijayarani S et al., [18] performed a comparative analysis of traditional association rule mining algorithms for data streams. The algorithms are APRIORI, APRIORI PT (Prefix Tree) and APRIORI MR (Map/Reduce). Different sizes of windows and threshold values are used and it concluded that APRIORI MR has produced good results than other algorithms.

Charu C. Aggarwal, [4] presented the detailed information about data streams. He discussed how to relate variant data mining technologies to data streams for supportive and unknown data extraction. He also discussed data stream clustering, data stream classification, association rule mining algorithm in data stream and frequent pattern mining.

Kuldeep Malik et al., [13] explained the FP-Growth algorithm and he proposed Enhanced FP-Growth Algorithm. He defined the Enhanced FP- Growth is working without prefix tree and any other complex data structure and he has proved that Enhanced FP-Growth has produced good results than FP-Growth.

III. PROPOSED ALGORITHM

The system architecture of this research work is represented in Figure 1.



Fig 1. System Architecture

A. Dataset

The connect data set is used in this work. This dataset is available in <u>http://fimi.ua.ac.be/data/connect.dat</u>. It has 67,558 instances and 48 attributes. From this 1K, 2K and 5K instances are used in this work. In data streams, we assume that the nonstop arrival of data is partitioned into many windows with permanent size, i.e. W_1 , W_2 , W_3 , W_n . In this work, we have created five windows W_1 , W_2 , W_3 , W_4 , W_5 with the data set size of 1K, 2K and 5K.

B. Association Rule Generation

- In order to generate association rules, two traditional data mining algorithms are used. They are,
- ✓ APriori MR Apriori Map/Reduce Algorithm.
- ✓ FP- Tree Frequent Pattern Tree Growth Algorithm.

C. APRIORI MR Algorithm

Apriori-Map/Reduce algorithm sprints on equivalent Map/Reduce framework. Candidate generation of Apriori Map/Reduce algorithm is prone (C_{k+1}) function is to eliminate the non-frequent item set C_{k+1} by reducing the non-frequent item sets cannot be a subset of frequent item sets. Table 1 corresponds to the APRIORI MR algorithm [12].

Table 1. Pseudo Code for APRIORI Map/Reduce Algorithm

- Step 1. Map transaction t in a data supply to all Map nodes
- Step 2. Each Map node can handle m
- Step 3. Now, can use Candidate Map Cm_1 = size of 1 is a frequent item set in the node m
- Step 4. Reduce and compute candidate generation of C₁ and L₁ with all Cm₁
- Step 5. C_1 = size one of frequent item sets;
- Step 6. Calculate the Min_Support = Num/ total items;
- Step 7. Size 1 of frequent item sets Min_Support is L₁
- Step 8. Loop begins, For $(k=1; L_k!=0; k++)$ do
- Step 9. Each mapped node m is represent by L_k . Such as, L_{mk}
- Step 10. Sort and remove the duplicate item sets
- Step 11. Can use, $C_{m(k+1)} = L_k \text{ join_sort } L_{mk}$;



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Step 12. Reduce methods to use the APRIORI Property to compute the C_{k+1} do

Step 13. Each map node m is increment the count of $L_{m(k+1)}$ candidates. That are supplied by transaction t

Step 14. End.

Step 15. Now, Can use reduce method to find the L_{k+1} with $L_{m(k+1)}$ and Min_Support.

Step 16. Min_Support of frequent item set generated by size of k+1 is L_{k+1} .

Step 17. End

Step 18. Return U_kL_k;

D. FP Tree Growth Algorithm

FP- Tree Growth used the divide and conquers tactics for creating the frequent item sets. FP-growth is primarily used for mining frequent item sets without candidate generation [13]. It consist of one root tags as null, a set of item prefix sub trees as the children of the root, and a frequent item subtile table. Every node in the item prefix sub tree consists of three fields: item-name, count and node link where item-name records which item the node characterized; count records the amount of transactions signified by the fraction of path reaching this node, the node links to the next node in the FP- tree. Each item in the subtile table consists of two fields, item name and head of node link, which positions to the first node in the FP-tree transports the item name. The pseudo code of mining on the FP - tree is portrayed in Table 2.

Window Size	Threshold	1000 Ds	2000 Ds	5000 Ds	10,000 Ds
		Rules			
W1	σ = 25, C = 55	18554	7473	7473	7473
W2		19588	7676	7667	7664
W3		7989	11112	7986	7982
W4		10154	10154	10143	10143
W5		9814	9814	9814	9810
W1	σ = 45, C = 55	3000	3090	3112	5553
W2		3003	3060	3045	3078
W3		3689	3652	3691	3634
W4		8465	8434	8490	9456
W5		5497	5493	5461	5449

Table 2. Pseudo Code for FP-Tree Growth Algorithm

- Step 1. Input: constructed FP-tree
- Step 2. Output: complete set of frequent patterns
- Step 3. Method: Call FP-growth (FP-tree, null).
- Step 4. Procedure:
- Step 5. FP-growth (Tree, α)
- Step 6.
- Step 7. if Tree contains a single path P then
- Step 8. For each combination do generate patterns $\beta^{\nu} \alpha$ with
- i. Support = min sup of nodes in β .
- Step 9. Else For each header a_i in the header of Tree do {
- Step 10. Generate pattern $\beta = a_i \cup \alpha$ with support = a_i support;
- Step 11. Construct β .s conditional pattern base and then β .s conditional

i. FP-tree Tree β

- Step 12. If Tree β = null {}
- Step 13. Then call FP-growth (Tree β , β)
- Step 14. }

IV. SIMULATION RESULTS

The section describes the experimental results of **APRIORI MR** and **FP-Tree Growth algorithms**. The work is implemented on Tanagra tool. Totally, five windows W_1 , W_2 , W_3 , W_4 and W_5 are used in this research work. There are



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four different sizes of datasets; 1K, 2K, 5K and 10K is tested and their results are obtained. Different threshold values are applied for analyzing the results. The performance factors used in this analysis are number of rules generated and execution time.

Window Size	Threshold	1000 Ds	2000 Ds	5000 Ds	10,000 Ds
		Rules			
W1	σ = 25, C = 55	18554	7473	7473	7473
W2		19588	7676	7667	7664
W3		7989	11112	7986	7982
W4		10154	10154	10143	10143
W5		9814	9814	9814	9810
W1	σ = 45, C = 55	3000	3090	3112	5553
W2		3003	3060	3045	3078
W3		3689	3652	3691	3634
W4		8465	8434	8490	9456
W5		5497	5493	5461	5449

Table 3.1 APRIORI MR Algorithm for Rule Generatio

Table 3.1 depicted the number of rules generated using APRIORI MR Algorithm. Two different thresholds values $\sigma = 45$, C = 55 are used.



Fig 3.1 Association Rule generation – APRIORI MR Algorithm

Figure 3.1 represented the rule generated by the APRIORI MR algorithm. From this result, it is observed that the minimum threshold value has generated more number of rules.



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Window Size	Threshold	1000 Ds	2000 Ds	5000 Ds	10,000 Ds
		Time(ms)			
W1	σ = 25, C = 55	2170	1539	3439	6633
W2		2784	1595	3539	6831
W3		9594	3230	3722	7051
W4		1180	2053	4527	8742
W 5		1171	2007	4442	8534
W1	σ = 45, C = 55	140	187	296	531
W2		156	203	265	577
W3		203	156	297	515
W4		281	281	609	850
W5		156	219	437	655

Table 3.2. Apriori MR Algorithm for Time Computation

Table 3.2 shows the execution time required for APRIORI MR algorithm. Two different thresholds values $\sigma = 45$, C = 55 are used.



Figure 3.2 represented the execution time taken by the APRIORI MR algorithm.



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Window Size	Threshold	1000 Ds	2000 Ds	5000 Ds	10,000 Ds
		Rules			
W1	$\sigma = 25,$ C = 55	65988	68345	83678	89712
W2		67250	67945	82444	89677
W3		65455	69896	84990	90012
W4		62390	65990	81266	86320
W5		61233	66435	85484	88531
W1	$\sigma = 45,$ C = 55	41435	45412	42265	49710
W2		44289	47892	47811	50101
W3		49342	43672	48234	49875
W4		43861	47239	50420	51676
W5		50905	46021	51346	49945

Table 3.3 FP- Tree Growth Algorithm for Rule Generation

Table 3.3 shows the number of rule generated using FP-Tree Growth algorithm. Two different thresholds values $\sigma = 45$, C = 55 are used.



Fig 3.3 Association Rule Generation – FP Tree Growth Algorithm

Figure 3.3 illustrated the number of rule generated using FP-Tree Growth algorithm.



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Window Size	Threshold	1000 Ds	2000 Ds	5000 Ds	10,000 Ds
		Time (ms)			
W1	σ = 25, C = 55	1448	1210	1310	1348
W2		1454	1221	1437	1340
W3		1455	1221	1441	1359
W4		1440	1446	1515	1567
W5		1337	1474	1243	1440
W1	σ = 45, C = 55	905	917	912	949
W2		915	917	918	944
W3		917	914	934	955
W4		921	919	946	967
W5		916	921	938	973

Table 3.4 FP-Tree Growth Algorithm for Time Computation

Table 3.4 depicted the execution time required for FP-Tree Growth algorithm. Two different thresholds values $\sigma = 45$, C = 55 are used.



Fig 3.4 Execution Time for FP Tree Growth Algorithm

Figure 3.4 shows the execution time for FP-Tree Growth algorithm.

V. CONCLUSION AND FUTURE WORK

This research work has compared the traditional association rule mining algorithms for generating association rules in data streams. Traditional association rule mining algorithm was experimented very less number of rules generation and execution time is very high. So, we compared with a new proposed algorithm like as FP Growth and APRIORI MR Algorithm. From the experimental results, it is observed that the performance of FP-Tree Growth algorithm is efficient than APRIORI MR Algorithm. In future work, new algorithms are to be developed in order to reduce the number of rules and execution time.



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BIOGRAPHY

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