Sequential Pattern Mining

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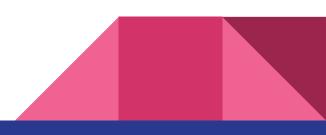
Introduction

Studies on Sequential Pattern Mining

- First introduced by **Agrawal** and **Srikant** in 1995
 - They presented three algorithms
 - AprioriAll
 - AprioriSome
 - DynamicSome
- Then in 1996 they presented **GSP** algorithm which was much faster than former algorithms and it also was generalized for more real life problems
- Pattern-growth methods: FreeSpan and PrefixSpan
- Mining closed sequential patterns: CloSpan

Applications

- Customer shopping sequences
 - First buy computer, then CD-ROM, and then digital camera, within 3 month.
- Medical treatments
- Natural disasters (e.g., earthquakes)
- Stocks and markets
- DNA sequences and gene structures



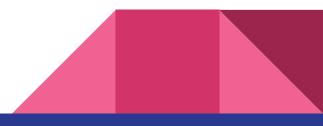
Problem Statement

We are given a database *D* of customer transactions. Each transaction consists of the following fields:

customer-id	transaction-time	items
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We want to find all **large sequences** that have a certain user-specified **minimum support.**

It is similar to the frequent itemsets mining, but with consideration of ordering.



Example of a database

Customer Id	Transaction Time Items Bough		
1	June 25 '93	30	
1	June 30 '93	90	
2	June 10 '93	10, 20	
2	June 15 '93	30	
2	June 20 '93	40, 60, 70	
3	June 25 '93	30, 50, 70	
4	June 25 '93	30	
4	June 30 '93	40, 70	
4	June 25 '93	90	
5	June 12 '93	90	

We convert DB to this form

Customer Id	Customer Sequence			
1	<(30)(90)>			
2	<(10 20) (30) (40 60 70)>			
3	<(30 50 70)>			
4	<(30) (40 70) (90)>			
5	<(90)>			

Definitions

Itemset and Sequence

An *itemset* is a non-empty set of items.

- We denote an itemset *i* by $(i_1 i_2 ... i_m)$

A Sequence is an ordered list of items.

- We denote a sequence s by $\langle s_1 s_2 \dots s_n \rangle$



Subsequence and supersequence

Given two sequences $\alpha = \langle a_1 a_2 \dots a_n \rangle$ and $\beta = \langle b_1 b_2 \dots b_m \rangle$:

- α is called a subsequence of β , if there exists integers $1 \le j_1 < j_2 < ... < j_n \le m$ such that $a_1 \subseteq b_1, a_2 \subseteq b_2, ..., a_n \subseteq b_{j_n}$
- β is called a supersequence of α

Example:

 $\alpha = <(a b) d > and \beta = <(a b c) (d e) >$



Apriori Property of Sequential Patterns

If a sequence *S* is not frequent, then none of the super-sequences of *S* is frequent.

Example: <h b> is infrequent -> so do <h a b> and <(a h) b>



Outline of the GSP Method

- Initially, every item in DB is a candidate of length-1
- For each level (i.e., sequences of length-k) do
 - Scan database to collect support count for each candidate sequence
 - Generate candidate length(k + 1) sequences from length-k frequent sequences using Apriori
 - Repeat until no frequent sequence or no candidate can be found



Major strength of **GSP** is its candidate pruning by Apriori property

Finding Length-1 Sequential Patterns

• Initial candidates:

o <a>, , <c>, <d>, <e>, <f>, <g>, <h>

• Scan database once, count support for candidates

 $min_sup = 2$

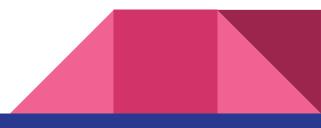
824.25	
Sequence	
<(bd)cb(ac)>	
<(bf)(ce)b(fg)>	
<(ah)(bf)abf>	
<(be)(ce)d>	
<a(bd)bcb(ade)></a(bd)bcb(ade)>	

Cand	Sup
<a>	3
	5
<c></c>	4
<d></d>	3
<e></e>	3
<f></f>	2
≤g≥	1
She	1

Generating Length-2 Candidates

	<a>		<c></c>	<d></d>	<e></e>	<f></f>
<a>	<aa></aa>	<ab></ab>	<ac></ac>	<ad></ad>	<ae></ae>	<af></af>
	<ba></ba>	<bb></bb>	<bc></bc>	<bd></bd>	<be></be>	<bf></bf>
<c></c>	<ca></ca>	<cb></cb>	<cc></cc>	<cd></cd>	<ce></ce>	<cf></cf>
<d></d>	<da></da>	<db></db>	<dc></dc>	<dd></dd>	<de></de>	<df></df>
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<f></f>	<fa></fa>	<fb></fb>	<fc></fc>	<fd></fd>	<fe></fe>	<ff></ff>

= 36



Generating Length-2 Candidates

	<a>		<c></c>	<d></d>	<e></e>	<f></f>
<a>		<(ab)>	<(ac)>	<(ad)>	<(ae)>	<(af)>
			<(bc)>	<(bd)>	<(be)>	<(bf)>
<c></c>				<(cd)>	<(ce)>	<(cf)>
<d></d>					<(de)>	<(df)>
<e></e>						<(ef)>
<f></f>						

= 15

Generating Length-2 Candidates

Using Apriori : 36 + 15 = **51** length-2 candidates

Without Apriori : (8 * 8) + (8 * 7) / 2 = 92 length- candidates

Apriori prunes 44.57% candidates

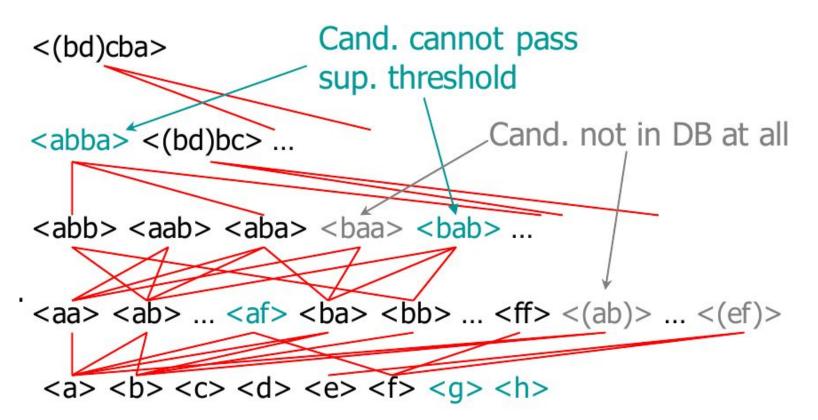


Finding Length-2 Sequential Patterns

- Scan database one more time, collect support count for each length-2 candidate
- There are 19 length-2 candidates which pass the minimum support threshold
 - They are length-2 sequential patterns



The GSP Mining Process



The GSP Algorithm

- Take sequences in form of <x> as length-1 candidates
- Scan database once, find F₁, the set of length-1 sequential patterns
- Let k = 1; while F_k is not empty do
 - Form C_{k+1} the set of length-(k + 1) candidates from F_k
 - If C_{k+1} is not empty, scan database once, find F_{k+1} , the set of length(k + 1) sequential patterns
 - Let k = k + 1

The Good, the Bad, and the Ugly



- *The Good*: benefits from the Apriori pruning which **reduces search space**
- *The Bad:* Scans the database multiple times
- *The Ugly:* Generates a **huge set of candidates** sequences



Why GSP is called Generalized Sequential Pattern Mining?

For practical use of **SPM**, in 1996 Agrawal and Srikant introduced three type of constraints that makes **SPM** problem more general and practical and since **GSP** support these constraints it is called **Generalized** sequential pattern mining.



Constraints that GSP Supports

Time Constraint

An ability for users to specify maximum and/or minimum time gaps between adjacent elements of the sequential pattern.



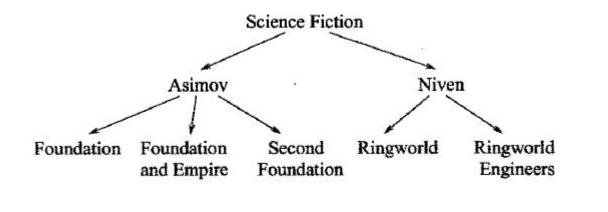
Sliding Window

That is, each element of the pattern can be contained in the union of the items bought in a set of transactions, as long as the difference between the maximum and minimum transaction-times is less than the size of a *sliding time window*.





An ability to define a taxonomy (*is-a* hierarchy) over the items in the data.





References

- R. Agrawal and R. Srikant. Mining Sequential Patterns. 1995
- R. Srikant and R. Agrawal. Mining Sequential Patterns.1996
- Jian Pei, Jiawei Han, Behzad Mortazavi-Asl, Helen Pinto, PrefixSpan: Mining Sequential Patterns Efficiently by Prefix-Projected Pattern Growth. 2001

