

# Web Data Mining

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## Example

Consider the following transactions:

(Bread, Eggs, Butter)

(Eggs, Bread, Milk)

(Cheese, Chips)

(Chips, Milk, Egg)

Find all rules that satisfy minimum support=0.5  
and minimum confidence=0.5







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# Road map

- Basic concepts of Association Rules
- Apriori algorithm
- Different data formats for mining
- Sequential pattern mining
- Summary

# Sequential pattern mining

- Association rule mining does not consider the order of transactions.
- In many applications such orderings are significant. E.g.,
  - in market basket analysis, it is interesting to know whether people buy some items in sequence,
    - e.g., buying bed first and then bed sheets some time later.
  - In Web usage mining, it is useful to find navigational patterns of users in a Web site from sequences of page visits of users

# Basic concepts

- Let  $I = \{i_1, i_2, \dots, i_m\}$  be a set of items.
- **Sequence**: An ordered list of itemsets.
- **Itemset/element**: A non-empty set of items  $X \subseteq I$ .  
We denote a sequence  $s$  by  $\langle a_1 a_2 \dots a_r \rangle$ , where  $a_i$  is an itemset, which is also called an **element** of  $s$ .
- An element (or an itemset) of a sequence is denoted by  $\{x_1, x_2, \dots, x_k\}$ , where  $x_j \in I$  is an item.
- We assume without loss of generality that items in an element of a sequence are in **lexicographic order**.



## Basic concepts (contd)

- **Size**: The **size** of a sequence is the number of elements (or itemsets) in the sequence.
- **Length**: The **length** of a sequence is the number of items in the sequence.
  - A sequence of length  $k$  is called  **$k$ -sequence**.
- A sequence  $s_1 = \langle a_1 a_2 \dots a_r \rangle$  is a **subsequence** of another sequence  $s_2 = \langle b_1 b_2 \dots b_v \rangle$ , or  $s_2$  is a **supersequence** of  $s_1$ , if there exist integers  $1 \leq j_1 < j_2 < \dots < j_{r-1} < j_r \leq v$  such that  $a_1 \subseteq b_{j_1}$ ,  $a_2 \subseteq b_{j_2}$ , ...,  $a_r \subseteq b_{j_r}$ . We also say that  $s_2$  **contains**  $s_1$ .

# An example

- Let  $I = \{1, 2, 3, 4, 5, 6, 7, 8, 9\}$ .
- Sequence  $\langle \{3\}\{4, 5\}\{8\} \rangle$  is **contained** in (or is a **subsequence** of)  $\langle \{6\} \{3, 7\}\{9\}\{4, 5, 8\}\{3, 8\} \rangle$ 
  - because  $\{3\} \subseteq \{3, 7\}$ ,  $\{4, 5\} \subseteq \{4, 5, 8\}$ , and  $\{8\} \subseteq \{3, 8\}$ .
  - However,  $\langle \{3\}\{8\} \rangle$  is not contained in  $\langle \{3, 8\} \rangle$  or vice versa.
  - The size of the sequence  $\langle \{3\}\{4, 5\}\{8\} \rangle$  is 3, and the length of the sequence is 4.

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# Objective

- Given a set  $S$  of **input data sequences** (or sequence database), the problem of mining sequential patterns is to find all the sequences that have **a user-specified minimum support**.
- Each such sequence is called a **frequent sequence**, or a **sequential pattern**.
- The **support** for a sequence is the fraction of total data sequences in  $S$  that contains this sequence.

# Example

**Table 1.** A set of transactions sorted by customer ID and transaction time

Customer ID	Transaction Time	Transaction (items bought)
1	July 20, 2005	30
1	July 25, 2005	90
2	July 9, 2005	10, 20
2	July 14, 2005	30
2	July 20, 2005	40, 60, 70
3	July 25, 2005	30, 50, 70
4	July 25, 2005	30
4	July 29, 2005	40, 70
4	August 2, 2005	90
5	July 12, 2005	90

## Example (cond)

**Table 2.** Data sequences produced from the transaction database in Table 1.

Customer ID	Data Sequence
1	$\langle\{30\} \{90\}\rangle$
2	$\langle\{10, 20\} \{30\} \{40, 60, 70\}\rangle$
3	$\langle\{30, 50, 70\}\rangle$
4	$\langle\{30\} \{40, 70\} \{90\}\rangle$
5	$\langle\{90\}\rangle$

**Table 3.** The final output sequential patterns

	Sequential Patterns with Support $\geq 25\%$
1-sequences	$\langle\{30\}\rangle, \langle\{40\}\rangle, \langle\{70\}\rangle, \langle\{90\}\rangle$
2-sequences	$\langle\{30\} \{40\}\rangle, \langle\{30\} \{70\}\rangle, \langle\{30\} \{90\}\rangle, \langle\{40, 70\}\rangle$
3-sequences	$\langle\{30\} \{40, 70\}\rangle$

# GSP mining algorithm

- Very similar to the Apriori algorithm

## Algorithm GSP( $S$ )

```
1   $C_1 \leftarrow \text{init-pass}(S);$  // the first pass over  $S$ 
2   $F_1 \leftarrow \{\langle \{f\} \rangle \mid f \in C_1, f.\text{count}/n \geq \text{minsup}\};$  //  $n$  is the number of sequences in  $S$ 
3  for ( $k = 2; F_{k-1} \neq \emptyset; k++$ ) do // subsequent passes over  $S$ 
4       $C_k \leftarrow \text{candidate-gen-SPM}(F_{k-1});$ 
5      for each data sequence  $s \in S$  do // scan the data once
6          for each candidate  $c \in C_k$  do
7              if  $c$  is contained in  $s$  then
8                   $c.\text{count}++;$  // increment the support count
9              end
10     end
11      $F_k \leftarrow \{c \in C_k \mid c.\text{count}/n \geq \text{minsup}\}$ 
12 end
13 return  $\bigcup_k F_k;$ 
```

**Fig. 12.** The GSP Algorithm for generating sequential patterns

# Candidate generation

**Function** candidate-gen-SPM( $F_{k-1}$ )

1. **Join step.** Candidate sequences are generated by joining  $F_{k-1}$  with  $F_{k-1}$ . A sequence  $s_1$  joins with  $s_2$  if the subsequence obtained by dropping the first item of  $s_1$  is the same as the subsequence obtained by dropping the last item of  $s_2$ . The candidate sequence generated by joining  $s_1$  with  $s_2$  is the sequence  $s_1$  extended with the last item in  $s_2$ . There are two cases:
  - the added item forms a separate element if it was a separate element in  $s_2$ , and is appended at the end of  $s_1$  in the merged sequence, and
  - the added item is part of the last element of  $s_1$  in the merged sequence otherwise.When joining  $F_1$  with  $F_1$ , we need to add the item in  $s_2$  both as part of an itemset and as a separate element. That is, joining  $\langle \{x\} \rangle$  with  $\langle \{y\} \rangle$  gives us both  $\langle \{x, y\} \rangle$  and  $\langle \{x\} \{y\} \rangle$ . Note that  $x$  and  $y$  in  $\{x, y\}$  are ordered.
2. **Prune step.** A candidate sequence is pruned if any one of its  $(k-1)$ -subsequence is infrequent (without minimum support).

**Fig. 13.** The candidate-gen-SPM() function

# An example

**Table 4.** Candidate generation: an example

Frequent 3-sequences	Candidate 4-sequences	
	after joining	after pruning
$\langle \{1, 2\} \{4\} \rangle$	$\langle \{1, 2\} \{4, 5\} \rangle$	$\langle \{1, 2\} \{4, 5\} \rangle$
$\langle \{1, 2\} \{5\} \rangle$	$\langle \{1, 2\} \{4\} \{6\} \rangle$	
$\langle \{1\} \{4, 5\} \rangle$		
$\langle \{1, 4\} \{6\} \rangle$		
$\langle \{2\} \{4, 5\} \rangle$		
$\langle \{2\} \{4\} \{6\} \rangle$		



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# Road map

- Basic concepts of Association Rules
- Apriori algorithm
- Different data formats for mining
- Mining with multiple minimum supports
- Mining class association rules
- Sequential pattern mining
- Summary

# Summary

- Association rule mining has been extensively studied in the data mining community.
- So is sequential pattern mining
- There are many efficient algorithms and model variations.
- Other related work includes
  - Multi-level or generalized rule mining
  - Constrained rule mining
  - Incremental rule mining
  - Maximal frequent itemset mining
  - Closed itemset mining
  - Rule interestingness and visualization
  - Parallel algorithms

# Example

Customer ID	Transaction Data	Item bought
1	1 <sup>st</sup> January	Bread, Jam
1	3 <sup>st</sup> January	Milk, Butter
2	2 <sup>nd</sup> January	Bread, Butter
2	5 <sup>th</sup> January	Milk
3	1 <sup>st</sup> January	Milk
3	4 <sup>th</sup> January	Butter, Jam
4	3 <sup>rd</sup> January	Bread, Butter, Jam

Find sequence with at least 50% support





