

SS-ZG548: ADVANCED DATA MINING

Lecture-14: Clustering on Data Stream, Big Data



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Sequence Data

- Sequence Data: $S = \langle e_1, e_2, e_3, \dots \rangle$ attributed with specific time
- Each element e_i is a list of events $\{i_1, i_2, \dots, i_k\}$
- Subsequence: $\langle a_1, a_2, \dots, a_n \rangle$ is contained in $\langle b_1, b_2, \dots, b_m \rangle$ if $\exists i_1 < i_2 < i_3 < \dots < i_n$ such that $a_1 \subseteq b_{i_1}, a_2 \subseteq b_{i_2}, \dots, a_n \subseteq b_{i_n}$
- Support for a sequence in database is the fraction that contains it

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Frequent sequence have support $\geq \text{minsup}$

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Frequent sequence have support $\geq \text{minsup}$

Consider sequences

$A = \langle \{1, 2, 4\}, \{2, 3\}, \{5\} \rangle$

$B = \langle \{1, 2\}, \{2, 3, 4\} \rangle$

$C = \langle \{1, 2\}, \{2, 3, 4\}, \{2, 4, 5\} \rangle$

$D = \langle \{2\}, \{3, 4\}, \{4, 5\} \rangle$

$E = \langle \{1, 3\}, \{2, 4, 5\} \rangle$

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Sequence	Support
$\langle \{1, 2\} \rangle$	

Sequence Data

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$E = \langle \{1, 3\}, \{2, 4, 5\} \rangle$

Sequence	Support
$\langle \{1, 2\} \rangle$	60%
$\langle \{2, 4\} \rangle$	80%
$\langle \{1\}, \{2\} \rangle$	80%
$\langle \{1, 2\}, \{2, 3\} \rangle$	60%

Generalized Sequential Pattern (GSP)¹

S-1 First pass to yield all 1-element frequent sequences

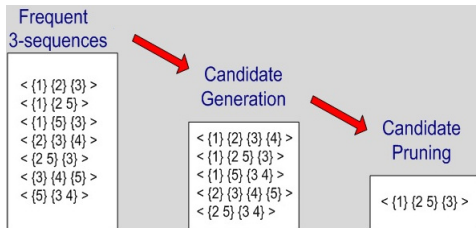
S-2 Repeat until new frequent sequences are found

- ▶ **Candidate Generation:** merge pairs found in $k - 1^{th}$ pass. w_1 and w_2 can be merged if subsequences obtained by removal of first element of w_1 and last element of w_2 are same
- ▶ **Candidate Pruning:** Prune candidates that contain a subsequence which is infrequent in $k - 1$ subsequences
- ▶ **Support Counting:** Need new pass to database
- ▶ **Candidate Elimination:** Involves thresholding based on minsup

$\langle \{1\}, \{2, 3\}\{4\} \rangle$ and $\langle \{2, 3\}, \{4, 5\} \rangle$	$\langle \{1\}, \{2, 3\}, \{4, 5\} \rangle$
$\langle \{1\}, \{2, 3\}\{4\} \rangle$ and $\langle \{2, 3\}, \{4\}, \{5\} \rangle$	$\langle \{1\}, \{2, 3\}, \{4\}, \{5\} \rangle$
$\langle \{1\}, \{2, 6\}\{4\} \rangle$ and $\langle \{1\}, \{2, 6\}, \{4\} \rangle$	Can not be merged

¹Generalized Sequential Pattern (GSP), Srikant and Agrawal, In EDBT 1996

Recap: Pruning in GSP



Candidate Generation

Frequent 3-sequences

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Candidate Generation

Frequent 3-sequences

$$\begin{aligned} & \langle \{1\} \{2\} \{3\} \rangle \\ & \langle \{1\} \{2\ 5\} \rangle \\ & \langle \{1\} \{5\} \{3\} \rangle \\ & \langle \{2\} \{3\} \{4\} \rangle \\ & \langle \{2\ 5\} \{3\} \rangle \\ & \langle \{3\} \{4\} \{5\} \rangle \\ & \langle \{5\} \{3\ 4\} \rangle \end{aligned}$$

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 ~~$\langle \{2\} \{3\} \{1\} \rangle$~~ ✓
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 ~~$\langle \{2\} \{3\} \{1\} \rangle$~~
 ~~$\langle \{2\} 5 \{3\} \rangle$~~ ✓
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 $\langle \{5\} \{3\} \{4\} \rangle$ ✓

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 ~~$\langle \{2\} 5 \{3\} \rangle$~~
 ~~$\langle \{3\} \{4\} \{5\} \rangle$~~ ✓
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 ~~$\langle \{5\} \{3 \ 1\} \rangle$~~ ✓

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Candidate Generation

$$\begin{aligned} & \langle \{1\} \{2\} \{3\} \{4\} \rangle \\ & \langle \{1\} \{2\ 5\} \{3\} \rangle \\ & \langle \{1\} \{5\} \{3\ 4\} \rangle \\ & \langle \{2\} \{3\} \{4\} \{5\} \rangle \\ & \langle \{2\ 5\} \{3\ 4\} \rangle \end{aligned}$$

Candidate Pruning

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Candidate Generation

< {1} {2} {3} {4} >
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Candidate Pruning

Candidate Generation

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Candidate Pruning

Candidate Generation

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Frequent 3-sequences

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Candidate Pruning

Candidate Generation

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Frequent 3-sequences

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Candidate Pruning

Candidate Generation

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$\langle \{2\} \{3\} \{4\} \rangle$ ✓
 $\langle \{1\} \{3\} \{4\} \rangle$ ✗
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Frequent 3-sequences

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Candidate Pruning

Candidate Generation

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Frequent 3-sequences

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Candidate Pruning

Candidate Generation

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Candidate Pruning

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Candidate Pruning

Candidate Generation

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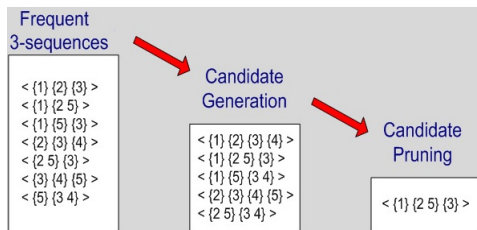
Frequent 3-sequences

$\langle \{1\} \{2\} \{3\} \rangle$
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Candidate Pruning

$\langle \{1\} \{2\} \{5\} \{3\} \rangle$

GSP: Candidate Generation



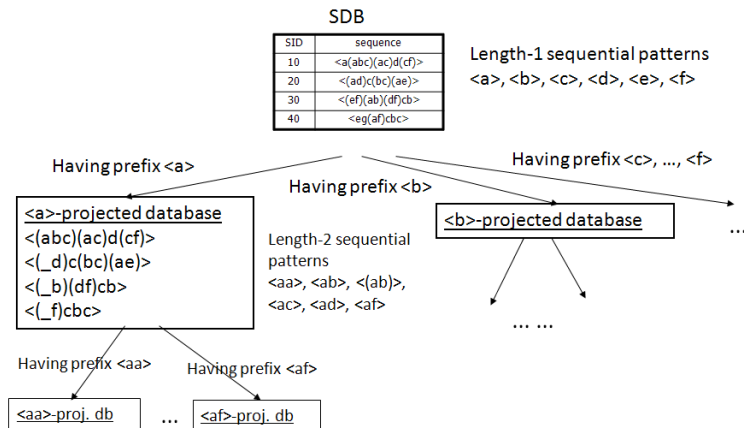
Issues:

- Huge number of candidate sets. n frequent 1-length candidate would generate $n^2 + \frac{n*(n-1)}{2}$ two-length candidate
- Multiple scans of the database
- Mining n -length sequential patterns need $\sum_{i=1}^n {}^nC_i = 2^n - 1$ number of short candidates. It is exponential

One can use prefix projections approach similar to FP-Growth

Pseudo-Projections²

When things can fit in main memory

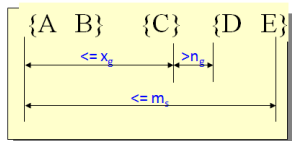


²Han, Jiawei and Pei, Jian and Mortazavi-Asl, Behzad and Pinto, Helen and Chen, Qiming and Dayal, Umeshwar and Hsu, MC, "Prefixspan: Mining sequential patterns efficiently by prefix-projected pattern growth" In proceedings of international conference on data engineering, pages 215–224, 2001

Application: Hotlink Assignment

- Hotlink are the mostly visited pages
- Website is modeled as graph, where pages are nodes and links are edges
- Web logs stores sequences of user clicks
- One sequence one session
- Intermediate pages could be navigational or target
- Use sequence mining to Mark start and end page of frequent sessions

Time Constraints



x_g : max-gap

n_g : min-gap

m_s : maximum span

$x_g = 2, n_g = 0, m_s = 4$

Data sequence	Subsequence	Contain?
$\langle \{2,4\} \{3,5,6\} \{4,7\} \{4,5\} \{8\} \rangle$	$\langle \{6\} \{5\} \rangle$	Yes
$\langle \{1\} \{2\} \{3\} \{4\} \{5\} \rangle$	$\langle \{1\} \{4\} \rangle$	No
$\langle \{1\} \{2,3\} \{3,4\} \{4,5\} \rangle$	$\langle \{2\} \{3\} \{5\} \rangle$	Yes
$\langle \{1,2\} \{3\} \{2,3\} \{3,4\} \{2,4\} \{4,5\} \rangle$	$\langle \{1,2\} \{5\} \rangle$	No

Approaches

- 1 Mine without timing constraint and post-process discovered patterns
- 2 Modify GSP to directly prune candidates violating timing

Anti Monotone Property

If a set is frequent all its subset must be frequent. If a set fails the test all its super set also must fail.

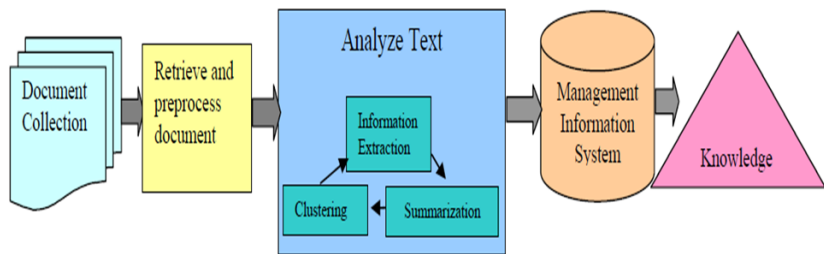
Consider sequences $A = \langle \{1, 2, 4\}, \{2, 3\}, \{5\} \rangle$
 $B = \langle \{1, 2\}, \{2, 3, 4\} \rangle$ $C = \langle \{1, 2\}, \{2, 3, 4\}, \{2, 4, 5\} \rangle$
 $D = \langle \{2\}, \{3, 4\}, \{4, 5\} \rangle$ $E = \langle \{1, 3\}, \{2, 4, 5\} \rangle$

- Let $x_g = 1(max - gap)$, $n_g = 0(min - gap)$, $m_s = 5(maxspan)$, $minSup = 60\%$
- What is support for $\langle \{2\}, \{5\} \rangle$? 40%
- What is support for $\langle \{2\}, \{3\}, \{5\} \rangle$? 60%

Anti Monotone Property does **not** holds, so these properties can not be pushed in GSP

Text Mining

- Computer are bad to handle slang, spelling variations, contextual meaning and unstructured data
- Text is less structured
- Applications involves 1) Information Extraction, 2) Topic Tracking, 3) Summarization, 4) Categorization, 5) Clustering, 6) Concept Linkage, 7) Information Visualization, 8) Question Answering, *etc.*
- Starts with 1) Identity keywords and phrases, 2) Relationship within text



Text Representation

- Binary term-document incidence matrix

	Antony and Cleopatra	Julius Caesar	The Tempest	Hamlet	Othello	Macbeth
Antony	1	1	0	0	0	1
Brutus	1	1	0	1	0	0
Caesar	1	1	0	1	1	1
Calpurnia	0	1	0	0	0	0
Cleopatra	1	0	0	0	0	0
mercy	1	0	1	1	1	1
worser	1	0	1	1	1	0

Document is represented as a binary vector $\in \{0, 1\}^{|V|}$

- Term-document count matrices

	Antony and Cleopatra	Julius Caesar	The Tempest	Hamlet	Othello	Macbeth
Antony	157	73	0	0	0	0
Brutus	4	157	0	1	0	0
Caesar	232	227	0	2	1	1
Calpurnia	0	10	0	0	0	0
Cleopatra	57	0	0	0	0	0
mercy	2	0	3	5	5	1
worser	2	0	1	1	1	0

Document is represented as a count vector $\in \mathbb{N}^{|V|}$

Text Representation

- Bag of words: order of words is not important. See “Jon is lighter than Bob” and “Bob is lighter than Jon”
- Term frequency ($tf_{t,d}$): is number of times the term t occurs in document d . Relevance may not increase proportionally with term frequency.
- Log-frequency weighting: $w_{t,d} = \log(tf_{t,d})$ if $tf_{t,d} > 0$ otherwise zero. Consider following matching score b/w two documents

$$score = \sum_{t \in D_1 \cap D_2} w_{t,D_1}$$

Score is zero if no term of query document D_2 is present in D_1 .

Text Representation

- **Document frequency:** Frequent terms are less informative than rare terms. df_t is number of documents that contain term t .
- **Inverse document frequency:** If we have N documents then $idf_t = \log(N/df_t)$
- **Collection frequency:** How many times term t appeared in all the document.
- **tfidf weighting:** $tfidf_{t,d} = \log(1 + tf_{t,d}) \times \log(N/df_t)$

$$score = \sum_{t \in q \cap d} tfidf_{t,d}$$

This is the most used method to determine similarity.

Text Representation

	Antony and Cleopatra	Julius Caesar	The Tempest	Hamlet	Othello	Macbeth
Antony	1	1	0	0	0	1
Brutus	1	1	0	1	0	0
Caesar	1	1	0	1	1	1
Calpurnia	0	1	0	0	0	0
Cleopatra	1	0	0	0	0	0
mercy	1	0	1	1	1	1
worser	1	0	1	1	1	0

	Antony and Cleopatra	Julius Caesar	The Tempest	Hamlet	Othello	Macbeth
Antony	157	73	0	0	0	0
Brutus	4	157	0	1	0	0
Caesar	232	227	0	2	1	1
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Cleopatra	57	0	0	0	0	0
mercy	2	0	3	5	5	1
worser	2	0	1	1	1	0

	Antony and Cleopatra	Julius Caesar	The Tempest	Hamlet	Othello	Macbeth
Antony	5.25	3.18	0	0	0	0.35
Brutus	1.21	6.1	0	1	0	0
Caesar	8.59	2.54	0	1.51	0.25	0
Calpurnia	0	1.54	0	0	0	0
Cleopatra	2.85	0	0	0	0	0
mercy	1.51	0	1.9	0.12	5.25	0.88
worser	1.37	0	0.11	4.15	0.25	1.95

- Generally dimensionality reduction is also required
- How document classification? use k-NN, Naive Bayes, SVM ...
- How document clustering? use k-Means, Hierarchical, Agglomerative

Statistics

There were 100 images in a box. 30 of them were containing lion. I asked Bob to separate all the pics of lion. He showed me 60 but, lion was not in 40 of them.

- True positives (TP): 20
- True negatives (TN): 30
- T1-Error: False positives (FP): 40
- T2-Error: False negatives (FN): 10

Confusion Matrix

		Experiment	
		T	F
Ground Truth	T	20	10
	F	40	30

Accuracy: $((20+30)/100)*100\%$,

Precision: $(20/60)*100\%$,

Recall (true positive rate or Sensitivity): $(20/(20+10))*100\%$,

Specificity (true negative rate): $(30/(40+30))*100\%$,

F Score: $(\text{Precision} + \text{Recall})/2$,

F1 Measure: Harmonic mean of Precision and Recall

Thank You!

Thank you very much for your attention!

Queries ?