

### Advanced Frequent Pattern Mining

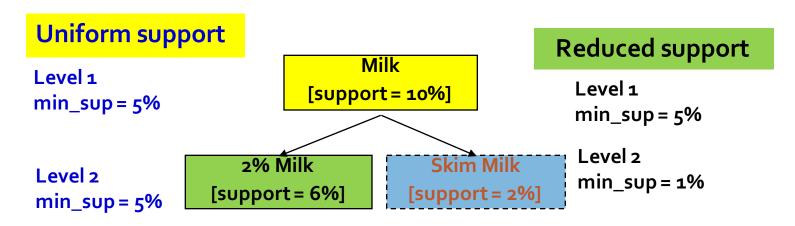
- Mining Diverse Patterns
- Constraint-Based Frequent Pattern Mining
- Mining High-Dimensional Data and Colossal Patterns
- Sequential Pattern Mining
- Graph Pattern Mining

### Mining Diverse Patterns

- Mining Multiple-Level Associations
- Mining Multi-Dimensional Associations
- Mining Quantitative Associations
- Mining Negative Correlations
- Mining Compressed Patterns

## Mining Multiple-Level Frequent Patterns

- Items often form hierarchies
  - Ex.: Dairyland 2% milk; Wonder wheat bread
- How to set min-support thresholds?
  - Uniform min-support across multiple levels (reasonable?)
  - Level-reduced min-support: Items at the lower level are expected to have lower support



### Redundancy Filtering at Mining Multi-Level Associations

- Multi-level association mining may generate many redundant rules
- Redundancy filtering: Some rules may be redundant due to "ancestor" relationships between items

(Suppose the 2% milk sold is about ¼ of milk sold in gallons)

- milk  $\Rightarrow$  wheat bread [support = 8%, confidence = 70%] (1)
- 2% milk  $\Rightarrow$  wheat bread [support = 2%, confidence = 72%] (2)
- A rule is redundant if its support is close to the "expected" value, according to its "ancestor" rule, and it has a similar confidence as its "ancestor"
  - Rule (1) is an ancestor of rule (2), which one to prune?

### Customized Min-Supports for Different Kinds of Items

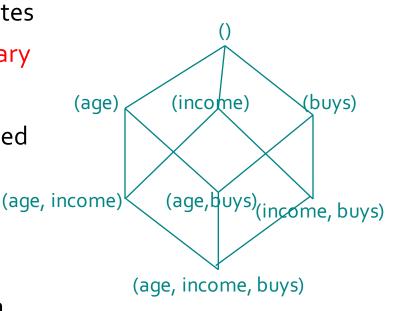
- We have used the same min-support threshold for all the items or item sets to be mined in each association mining
- In reality, some items (e.g., diamond, watch, ...) are valuable but less frequent
- It is necessary to have customized min-support settings for different kinds of items
- One Method: Use group-based "individualized" min-support
  - E.g., {diamond, watch}: 0.05%; {bread, milk}: 5%; ...

### Mining Multi-Dimensional Associations

- Single-dimensional rules (e.g., items are all in "product" dimension)
  - buys(X, "milk")  $\Rightarrow$  buys(X, "bread")
- Multi-dimensional rules (i.e., items in ≥ 2 dimensions or predicates)
  - Inter-dimension association rules (no repeated predicates)
    - age(X, "18-25") ∧ occupation(X, "student") ⇒ buys(X, "coke")
  - Hybrid-dimension association rules (repeated predicates)
    - age(X, "18-25")  $\land$  buys(X, "popcorn")  $\Rightarrow$  buys(X, "coke")
- Attributes can be categorical or numerical
  - Categorical Attributes (e.g., profession, product: no ordering among values): Data cube for inter-dimension association
  - Quantitative Attributes: Numeric, implicit ordering among values—discretization, clustering, and gradient approaches

## Mining Quantitative Associations

- Mining associations with numerical attributes
  - Ex.: Numerical attributes: age and salary
- Methods
  - Static discretization based on predefined concept hierarchies
    - Data cube-based aggregation
  - Dynamic discretization based on data distribution
  - Clustering: Distance-based association
    - First one-dimensional clustering, then association
  - Deviation analysis:
    - Gender = female ⇒ Wage: mean=\$7/hr (overall mean = \$9)



# Mining Extraordinary Phenomena in Quantitative Association Mining

- Mining extraordinary (i.e., interesting) phenomena
  - Ex.: Gender = female  $\Rightarrow$  Wage: mean=\$7/hr (overall mean = \$9)
  - LHS: a subset of the population
  - RHS: an extraordinary behavior of this subset
- The rule is accepted only if a statistical test (e.g., Z-test) confirms the inference with high confidence
- Subrule: Highlights the extraordinary behavior of a subset of the population of the super rule
  - Ex.: (Gender = female)  $^{\land}$  (South = yes)  $\Rightarrow$  mean wage = \$6.3/hr
- Rule condition can be categorical or numerical (quantitative rules)
  - Ex.: Education in [14-18] (yrs)  $\Rightarrow$  mean wage = \$11.64/hr
- Efficient methods have been developed for mining such extraordinary rules (e.g., Aumann and Lindell@KDD'99)

#### Rare Patterns vs. Negative Patterns

- Rare patterns
  - Very low support but interesting (e.g., buying Rolex watches)
  - How to mine them? Setting individualized, group-based minsupport thresholds for different groups of items
- Negative patterns
  - Negatively correlated: Unlikely to happen together
  - Ex.: Since it is unlikely that the same customer buys both a
     Ford Expedition (an SUV car) and a Ford Fusion (a hybrid car),
     buying a Ford Expedition and buying a Ford Fusion are likely
     negatively correlated patterns
  - How to define negative patterns?

#### Defining Negative Correlated Patterns

- A support-based definition
  - If itemsets A and B are both frequent but rarely occur together,
     i.e., sup(A U B) << sup(A) × sup(B)</li>
  - Then A and B are negatively correlated
- the definition of *lift*?
- Is this a good definition for large transaction datasets?
- Ex.: Suppose a store sold two needle packages A and B 100 times each, but only one transaction contained both A and B
  - When there are in total 200 transactions, we have
    - $s(A \cup B) = 0.005$ ,  $s(A) \times s(B) = 0.25$ ,  $s(A \cup B) << s(A) \times s(B)$
  - But when there are 10<sup>5</sup> transactions, we have
    - $s(A \cup B) = 1/10^5$ ,  $s(A) \times s(B) = 1/10^3 \times 1/10^3$ ,  $s(A \cup B) > s(A) \times s(B)$
  - What is the problem? Null transactions: The support-based definition is not null-invariant!

## Defining Negative Correlation: Need Null-Invariance in Definition

- A good definition on negative correlation should take care of the null-invariance problem
  - Whether two itemsets A and B are negatively correlated should not be influenced by the number of nulltransactions
- A Kulczynski measure-based definition
  - If itemsets A and B are frequent but  $(P(A|B) + P(B|A))/2 < \epsilon$ , where  $\epsilon$  is a negative pattern threshold, then A and B are negatively correlated
- For the same needle package problem:
  - No matter there are in total 200 or 105 transactions
  - If  $\epsilon$  = 0.01, we have  $(P(A|B) + P(B|A))/2 = (0.01 + 0.01)/2 < \epsilon$

## Mining Compressed Patterns

- Why mining compressed patterns?
  - Too many scattered patterns but not so meaningful
- Pattern distance measure

$$Dist(P_1, P_2) = 1 - \frac{|T(P_1) \cap T(P_2)|}{|T(P_1) \cup T(P_2)|}$$

- $\delta$ -clustering: For each pattern P, find all  $\frac{P}{\Gamma}$  patterns which can be expressed by P and whose distance to P is within  $\delta$  ( $\delta$ -cover)
- All patterns in the cluster can be represented by P
- Method for efficient, direct mining of compressed frequent patterns (e.g., D. Xin, J. Han, X. Yan, H. Cheng, "On Compressing Frequent Patterns", Knowledge and Data Engineering, 60:5-29, 2007)

Pat-ID	Item-Sets	Support
P1	{38,16,18,12}	205227
P <sub>2</sub>	{38,16,18,12,17}	205211
P <sub>3</sub>	{39,38,16,18,12,17}	101758
P <sub>4</sub>	{39,16,18,12,17}	161563
P <sub>5</sub>	{39,16,18,12}	161576

- Closed patterns
  - P1, P2, P3, P4, P5
  - Emphasizes too much on support
  - ☐ There is no compression
- Max-patterns
  - □ P3: information loss
- Desired output (a good balance):
  - □ P2, P3, P4

#### Advanced Frequent Pattern Mining

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## Why Constraint-Based Mining?

- Finding all the patterns in a dataset autonomously?— unrealistic!
  - Too many patterns but not necessarily user-interested!
- Pattern mining should be an interactive process
  - User directs what to be mined using a data mining query language (or a graphical user interface)
- Constraint-based mining
  - User flexibility: provides constraints on what to be mined
  - Optimization: explores such constraints for efficient mining
    - Constraint-based mining: Constraint-pushing, similar to push selection first in DB query processing

## Constraints in General Data Mining

A data mining query can be in the form of a metarule or with the following language primitives.

- Knowledge type constraint:
  - Ex.: classification, association, clustering, outlier finding, ....
- Data constraint using SQL-like queries
  - Ex.: find products sold together in NY stores this year
- Dimension/level constraint
  - Ex.: in relevance to region, price, brand, customer category
- Rule (or pattern) constraint
  - Ex.: small sales (price < \$10) triggers big sales (sum > \$200)
- Interestingness constraint
  - Ex.: strong rules: min\_sup ≥ 0.02, min\_conf ≥ 0.6, min\_correlation ≥ 0.7

## Meta-Rule Guided Mining

A meta-rule can contain partially instantiated predicates & constants

```
- P_1(X, Y) \wedge P_2(X, W) \Rightarrow buys(X, "iPad")
```

• The resulting mined rule can be

- age(X, "15-25") 
$$^{\circ}$$
 profession(X, "student")  $\Rightarrow$  buys(X, "iPad")

• In general, (meta) rules can be in the form of

$$-P_1 \wedge P_2 \wedge ... \wedge P_1 \Rightarrow Q_1 \wedge Q_2 \wedge ... \wedge Q_r$$

- Method to find meta-rules
  - Find frequent (I + r) predicates (based on min-support)
  - Push constants deeply when possible into the mining process
  - Also, push min\_conf, min\_correlation, and other measures as early as possible (measures acting as constraints)

## Different Kinds of Constraints Lead to Different Pruning Strategies

- Constraints can be categorized as
  - Pattern space pruning constraints vs. data space pruning constraints
- Pattern space pruning constraints
  - Anti-monotonic: If constraint c is violated, its further mining can be terminated (=no superset)
  - Monotonic: If c is satisfied, no need to check c again (=all supersets)
  - Succinct: if the constraint c can be enforced by directly manipulating the data (e.g., data pruning)
  - Convertible: c can be converted to monotonic or anti-monotonic if items can be properly ordered in processing
- Data space pruning constraints
  - Data succinct: Data space can be pruned at the initial pattern mining process
  - Data anti-monotonic: If a transaction t does not satisfy c, then t can be pruned to reduce data processing effort

# Pattern Space Pruning with Pattern Anti-Monotonicity

- Constraint c is anti-monotone
  - If an itemset S violates constraint c, so does any of its superset
  - That is, mining on itemset S can be terminated
- Ex. 1: c1: sum(S.price) ≤ v is antimonotone
- Ex. 2: c2: range(S.profit) ≤ 15 is antimonotone
  - Itemset ab violates c2 (range(ab) = 40)
  - So does every superset of ab
- Ex. 3. c3: sum(S.Price) ≥ v is not antimonotone
- Ex. 4. Is c4: support(S) ≥ σ antimonotone?
  - Yes! Apriori pruning is essentially pruning with an anti-monotonic constraint!

TID	Transaction	
10	a, b, c, d, f, h	
20	b, c, d, f, g, h	
30	b, c, d, f, g	
40	a, c, e, f, g	
min_sup = 2		
price(item)>o		

ltem	Profit
a	40
b	0
С	-20
d	-15
е	-30
f	-10
g	20
h	5

## Pattern Monotonicity and Its Roles

- A constraint c is monotone: if an itemset S satisfies the constraint c, so does any of its superset
  - That is, we do not need to check c in subsequent mining
- Ex. 1: c1: sum(S.Price) ≥ v is monotone
- Ex. 2: c2: min(S.Price) ≤ v is monotone
- Ex. 3: c3: range(S.profit) ≥ 15 is monotone
  - Itemset ab satisfies c3
  - So does every superset of ab

TID	Transaction		
10	a, b, c, d, f, h		
20	b, c, d, f, g, h		
30	b, c, d, f, g		
40	a, c, e, f, g		
min_sup = 2			
price(item)>o			

ltem	Profit
a	40
b	0
С	-20
d	-15
е	-30
f	-10
g	20
h	5

## Data Space Pruning with Data Anti-Monotonicity

- A constraint c is data anti-monotone: In the mining process, if a data entry t cannot satisfy a pattern p under c, t cannot satisfy p's superset either
  - Data space pruning: Data entry t can be pruned
- Ex. 1:  $c_1$ :  $sum(S.Profit) \ge v$  is data antimonotone
  - Let constraint  $c_1$  be: sum{S.Profit} ≥ 25
    - T<sub>30</sub>: {b, c, d, f, g} can be removed since none of their combinations can make an S whose sum of the profit is ≥ 25
- Ex. 2:  $c_2$ :  $min(S.Price) \le v$  is data antimonotone
  - Consider v = 5 but every item in transaction  $T_{50}$  has a price higher than 10
- Ex. 3: c<sub>3</sub>: range(S.Profit) ≥ 25 is data antimonotone

TID	Transaction		
10	a, b, c, d, f, h		
20	b, c, d, f, g, h		
30	b, c, d, f, g		
40	a, c, e, f, g		
min_sup = 2			
price(item)>o			

Item	Profit
а	40
b	0
С	-20
d	-15
е	-30
f	-10
g	20
h	5

# Succinctness: Pruning Both Data and Pattern Spaces

- Succinctness: if the constraint c can be enforced by directly manipulating the data
- Ex. 1: To find those patterns without item i
  - Remove i from DB and then mine (pattern space pruning)
- Ex. 2: To find those patterns containing item i
  - Mine only i-projected DB (data space pruning)
- Ex. 3:  $c_3$ : min(S.Price)  $\leq v$  is succinct
  - Start with only items whose price ≤ v (pattern space pruning) and remove transactions with high-price items only (data space pruning)
- Ex. 4:  $c_4$ : sum(S.Price)  $\geq v$  is not succinct
  - It cannot be determined beforehand since sum of the price of itemset S keeps increasing

## Convertible Constraints: Ordering Data in Transactions

- Convert tough constraints into (anti-) monotone by proper ordering of items in transactions
- Examine c1: avg(S.profit) > 20
  - Order items in value-descending order
    - <a, g, f, h, b, d, c, e>
  - An itemset ab violates c1 (avg(ab) = 20)
    - So does ab\* (i.e., ab-projected DB)
    - C1: anti-monotone if patterns grow in the right order!
- Can item-reordering work for Apriori?
  - Does not work for level-wise candidate generation!
  - avg(agf) = 23.3 > 20, but avg(gf) = 15 < 20

		ltem	Profit
		a	40
		b	0
		С	-20
		d	-15
		е	-30
		f	10
min_sup = 2		g	20
price(item)>o		h	5
TID	Transaction		

a, b, c, d, f, h

b, c, d, f, g, h

b, c, d, f, g

a, c, e, f, g

10

20

30

40

#### How to Handle Multiple Constraints?

- It is beneficial to use multiple constraints in pattern mining
- But different constraints may require potentially conflicting item-ordering
  - If there exists an order R making both c1 and c2 convertible, try to sort items in the order that benefits pruning most
  - If there exists conflict ordering between c1 and c2
    - Try to sort data and enforce one constraint first (which one?)
    - Then enforce the other when mining the projected databases
- Ex.  $c_1$ : avg(S.profit) > 20, and  $c_2$ : avg(S.price) < 50
  - Sorted in profit descending order and use c1 first (assuming  $c_1$  has more pruning power)
  - For each project DB, sort trans. in price ascending order and use c<sub>2</sub> at mining

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## Mining Long Patterns: Challenges

- Mining long patterns is needed in bioinformatics, social network analysis, software engineering, ...
  - But the methods introduced so far mine only short patterns (e.g., length < 10)</li>
- Challenges of mining long patterns
  - The curse of "downward closure" property of frequent patterns
    - Any sub-pattern of a frequent pattern is frequent
    - If  $\{a_1, a_2, ..., a_{100}\}$  is frequent, then  $\{a_1\}, \{a_2\}, ..., \{a_{100}\}, \{a_1, a_2\}, \{a_1, a_3\}, ..., \{a_{100}\}, \{a_1, a_2, a_3\}, ...$  are all frequent! There are about  $2^{100}$  such frequent itemsets!
  - No matter searching in breadth-first (e.g., Apriori) or depth-first (e.g., FPgrowth), if we still adopt the "small to large" step-by-step growing paradigm, we have to examine so many patterns, which leads to combinatorial explosion!

# Colossal Patterns: A Motivating Example

- Let min-support  $\sigma$ = 20 • # of closed/maximal patterns of size 20: about  $\binom{40}{20}$
- But there is only one pattern with size close to 40 (i.e., long or colossal)
  - $-\alpha = \{41, 42, ..., 79\}$  of size 39
- Q: How to find it without generating an exponential number of size-20 patterns?

```
T<sub>1</sub> = 2 3 4 ..... 39 40
T<sub>2</sub> = 1 3 4 ..... 39 40
: .
```

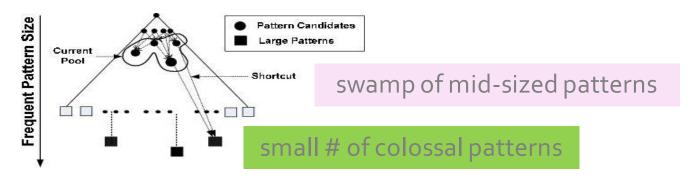
T60= 41 42 43 ... 79

The existing fastest mining algorithms (e.g., FPClose, LCM) fail to complete running

A new algorithm, *Pattern-Fusion*, outputs this colossal pattern in seconds

#### What Is Pattern-Fusion?

- Not strive for completeness (why?)
- Jump out of the swamp of the mid-sized intermediate "results"
- Strive for mining almost complete and representative colossal patterns: identify "short-cuts" and take "leaps"
- Key observation
  - The larger the pattern or the more distinct the pattern, the greater chance it will be generated from small ones
- Philosophy: Collection of small patterns hints at the larger patterns
- Pattern fusion strategy ("not crawl but jump"): Fuse small patterns together in one step to generate new pattern candidates of significant sizes



## Observation: Colossal Patterns and Core Patterns

- Suppose dataset D contains 4 colossal patterns (below) plus many small patterns
  - $\{a_1, a_2, ..., a_{50}\}$ : 40,  $\{a_3, a_6, ..., a_{99}\}$ : 60,  $\{a_5, a_{10}, ..., a_{95}\}$ : 80,  $\{a_{10}, a_{20}, ..., a_{100}\}$ : 100
- If you check the pattern pool of size-3, you may likely find
  - {a<sub>2</sub>, a<sub>4</sub>, a<sub>45</sub>}: ~40; {a<sub>3</sub>, a<sub>34</sub>, a<sub>39</sub>}: ~40; ..., {a<sub>5</sub>, a<sub>15</sub>, a<sub>85</sub>}: ~80, ..., {a<sub>20</sub>, a<sub>40</sub>, a<sub>85</sub>}: ~80, ...
- If you merge the patterns with similar support, you may obtain candidates of much bigger size and easily validate whether they are true patterns
- Core patterns of a colossal pattern  $\alpha$ : A set of subpatterns of  $\alpha$  that cluster around  $\alpha$  by sharing a similar support
- A colossal pattern has far more core patterns than a small-sized pattern
- A random draw from a complete set of pattern of size c would be more likely to pick a core pattern (or its descendant) of a colossal pattern
- A colossal pattern can be generated by merging a set of core patterns

#### Robustness of Colossal Patterns

• Core Patterns: For a frequent pattern  $\alpha$ , a subpattern  $\beta$  is a  $\tau$ -core pattern of  $\alpha$  if  $\beta$  shares a similar support set with  $\alpha$ , i.e.,

 $\frac{|D_{\alpha}|}{|D_{\beta}|} \ge \tau \quad 0 < \tau \le 1$  where  $\tau$  is called the core ratio

- (d,  $\tau$ )-robustness: A pattern  $\alpha$  is (d,  $\tau$ )-robust if d is the maximum number of items that can be removed from  $\alpha$  for the resulting pattern to remain a  $\tau$ -core pattern of  $\alpha$
- For a (d,  $\tau$ )-robust pattern  $\alpha$ , it has  $\Omega(2^d)$  core patterns
- Robustness of Colossal Patterns: A colossal pattern tends to have much more core patterns than small patterns
- Such core patterns can be clustered together to form "dense balls" based on pattern distance defined by

A random draw in the pattern space will hit somewhere in the ball with high probability

$$Dist(\alpha, \beta) = 1 - \frac{\left| D_{\alpha} \cap D_{\beta} \right|}{\left| D_{\alpha} \cup D_{\beta} \right|}$$

## The Pattern-Fusion Algorithm

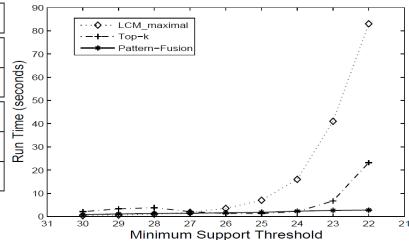
- Initialization (Creating initial pool): Use an existing algorithm to mine all frequent patterns up to a small size, e.g., 3
- Iteration (Iterative Pattern Fusion):
  - At each iteration, K seed patterns are randomly picked from the current pattern pool
  - For each seed pattern thus picked, we find all the patterns within a bounding ball centered at the seed pattern
  - All these patterns found are fused together to generate a set of super-patterns
  - All the super-patterns thus generated form a new pool for the next iteration
- Termination: when the current pool contains no more than K patterns at the beginning of an iteration

#### Experimental Results on Data Set: ALL

- ALL: A popular gene expression clinical data set on ALL-AML leukemia, with 38 transactions, each with 866 columns. There are 1,736 items in total.
  - When minimum support is high (e.g., 30), Pattern-Fusion gets all the largest colossal patterns with size greater than 85

Pattern Size	110	107	102	91	86	84	83
The complete set	1	1	1	1	1	2	6
Pattern-Fusion	1	1	1	1	1	1	4
Pattern Size	82	77	76	75	74	73	71
The complete set	1	2	1	1	1	2	1
Pattern-Fusion	0	2	0	1	1	1	1

Mining colossal patterns on a Leukemia dataset



Algorithm runtime comparison on another dataset

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# Sequence Databases and Sequential Patterns

- Sequential pattern mining has broad applications
  - Customer shopping sequences
    - Purchase a laptop first, then a digital camera, and then a smartphone, within 6 months
  - Medical treatments, natural disasters (e.g., earthquakes),
     science & engineering processes, stocks and markets, ...
  - Weblog click streams, calling patterns, ...
  - Software engineering: Program execution sequences, ...
  - Biological sequences: DNA, protein, ...
- Transaction DB, sequence DB vs. time-series DB
- Gapped vs. non-gapped sequential patterns
  - Shopping sequences, clicking streams vs. biological sequences

# Sequential Pattern and Sequential Pattern Mining

 Sequential pattern mining: Given a set of sequences, find the complete set of frequent subsequences (i.e., satisfying the min\_sup threshold)

A <u>sequence database</u>		Sequence
		<a(<u>abc)(a<u>c</u>)d(cf)&gt;</a(<u>
	20	<(ad)c(bc)(ae)>
	30	<(ef)( <u>ab</u> )(df) <u>c</u> b>
A <u>sequence:</u> < (ef) (ab) (df) c ,b >	40	<eg(af)cbc></eg(af)cbc>

- An element may contain a set of items (also called events)
- Items within an element are unordered and we list them alphabetically <a(bc)dc> is a <u>subsequence</u> of <<u>a(bc)d(cf)></u>
- Given support threshold min\_sup = 2, <(ab)c> is a sequential pattern

#### Sequential Pattern Mining Algorithms

- Algorithm requirement: Efficient, scalable, finding complete set, incorporating various kinds of user-specific constraints
- The Apriori property still holds: If a subsequence s<sub>1</sub> is infrequent, none of s<sub>1</sub>'s super-sequences can be frequent
- Representative algorithms
  - GSP (Generalized Sequential Patterns): Srikant & Agrawal @
     EDBT'96)
  - Vertical format-based mining: SPADE (Zaki@Machine Leanining'oo)
  - Pattern-growth methods: PrefixSpan (Pei, et al. @TKDE'04)
- Mining closed sequential patterns: CloSpan (Yan, et al. @SDM'o3)
- Constraint-based sequential pattern mining

#### GSP: Apriori-Based Sequential Pattern

#### Mining

- Initial candidates: All singleton sequences
  - <a>, <b>, <c>, <d>, <e>, <f>, <g>, <h>
- Scan DB once, count support for each candidate
- Generate length-2 candidate sequences

SID	Sequence
10	<(bd)cb(ac)>
20	<(bf)(ce)b(fg)>
30	<(ah)(bf)abf>
40	<(be)(ce)d>
50	<a(bd)bcb(ade)></a(bd)bcb(ade)>

min_sup = 2		
Cand.	sup	
<a></a>	3	
<b></b>	5	
<c></c>	4	
<d></d>	3	
<e></e>	3	
<f></f>	2	
<g></g>	1	
<h></h>	1	

	_					
	<a></a>	<b></b>	<c></c>	<d></d>	<e></e>	<f></f>
<a></a>	<aa></aa>	<ab></ab>	<ac></ac>	<ad></ad>	<ae></ae>	<af></af>
<b></b>	<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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	<a></a>	<b></b>	<c></c>	<d></d>	<e></e>	<f></f>
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<b></b>			<(bc)>	<(bd)>	<(be)>	<(bf)>
<c></c>				<(cd)>	<(ce)>	<(cf)>
<d></d>					<(de)>	<(df)>
<e></e>						<(ef)>
٠٤.						

Length-2 candidates: 36 + 15 = 51 Without Apriori pruning: 8\*8+8\*7/2=9 2 candidates

#### (Generalized Sequential

**GSP** 

Patterns): Srikant & Agrawal @ EDBT'96)

### **GSP Mining and Pruning**

- Repeat (for each level (i.e., length-k))
  - Scan DB to find length-k frequent sequences
  - Generate length-(k+1) candidate sequences from length-k
     frequent sequences using Apriori
  - set k = k+1
- Until no frequent sequence or no candidate can be found

# Sequential Pattern Mining in Vertical Data Format: The SPADE Algorithm

A sequence database is mapped to: <SID, EID>

Grow the subsequences (patterns) one item at a time by Apriori candidate

generation

SID	Sequence
1	<a(<u>abc)(a<u>c</u>)d(cf)&gt;</a(<u>
2	<(ad)c(bc)(ae)>
3	<(ef)( <u>ab</u> )(df) <u>c</u> b>
4	<eg(af)cbc></eg(af)cbc>

 $min\_sup = 2$ 

Ref: SPADE (<u>S</u>equential <u>PA</u>ttern <u>D</u>iscovery using <u>E</u>quivalent Class)
[M. Zaki 2001]

SID	EID	Items
1	1	a
1	2	abc
1	3	ac
1	4	d
1	5	$\operatorname{cf}$
2	1	ad
2	2	$\mathbf{c}$
2	3	$_{\mathrm{bc}}$
2	4	ae
3	1	$\operatorname{ef}$
3	2	ab
3	3	$\mathrm{d}\mathrm{f}$
3	4	$\mathbf{c}$
3	5	b
4	1	$\mathbf{e}$
4	2	g
4	3	af
4	4	$\mathbf{c}$
4	5	b
4	6	$\mathbf{c}$

	$\mathbf{a}$	1	О	
SID	EID	$\operatorname{SID}$	EID	
1	1	1	2	
1	2	2	3	
1	3	3	2	
2	1	3	5	
2	4	4	5	
3	2			
4	3			
	ab		ba	

	ab			ba		
SID	EID (a)	EID(b)	$\operatorname{SID}$	EID (b)	EID(a)	
1	1	$^2$	1	$^2$	3	
2	1	3	2	3	4	
3	2	5				
4	3	5				

	ŧ	aba		
SID	EID (a)	EID(b)	EID(a)	
1	1	2	3	
2	1	3	4	

# PrefixSpan: A Pattern-Growth Approach

- Prefix and suffix
  - Given <a(abc)(ac)d(cf)>
  - Prefixes: <a>, <aa>, <a(ab)>, <a(abc)>, ...
  - Suffix: Prefixes-based projection
- PrefixSpan Mining: Prefix Projections
  - Step 1: Find length-1 sequential patterns
    - <a>, <b>, <c>, <d>, <e>, <f>
  - Step 2: Divide search space and mine each projected DB
    - <a>-projected DB,
    - <b>-projected DB,
    - ...
    - <f>-projected DB, ...

SID	Sequence
10	<a(<u>abc)(a<u>c</u>)d(cf)&gt;</a(<u>
20	<(ad)c(bc)(ae)>
30	<(ef)( <u>ab</u> )(df) <u>c</u> b>
40	<eg(af)cbc></eg(af)cbc>

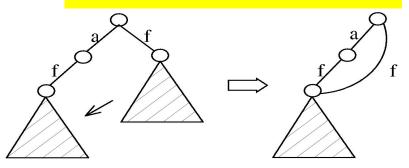
Prefix	Suffix (Projection)
<a></a>	<(abc)(ac)d(cf)>
<aa></aa>	<(_bc)(ac)d(cf)>
<ab></ab>	<(_c)(ac)d(cf)>

PrefixSpan (Prefix-projected Sequential pattern mining) Pei, et al. @TKDE'04

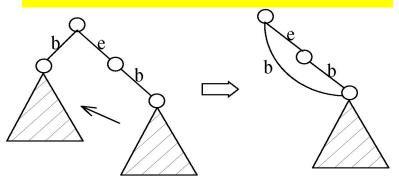
## CloSpan: Mining Closed Sequential Patterns

- A closed sequential pattern s: There exists no superpattern s' such that s' > s, and s' and s have the same support
- Which ones are closed? <abc>: 20,
   <abcd>: 20,
   <abcd>: 15
- Why directly mine closed sequential patterns?
  - Reduce # of (redundant) patterns
  - Attain the same expressive power
- Property  $P_1$ : If  $s \supset s_1$ , s is closed iff two project DBs have the same size
- Explore Backward Subpattern and Backward Superpattern pruning to prune redundant search space
- Greatly enhances efficiency (Yan, et al., SDM'o<sub>3</sub>)

Backward subpattern pruning



Backward superpattern pruning



# Constraint-Based Sequential-Pattern Mining

- Share many similarities with constraint-based itemset mining
- Anti-monotonic: If S violates c, the super-sequences of S also violate c
  - sum(S.price) < 150; min(S.value) > 10
- Monotonic: If S satisfies c, the super-sequences of S also do so
  - element\_count(S) > 5;  $S \supseteq \{PC, digital\_camera\}$
- Data anti-monotonic: If a sequence  $s_1$  with respect to S violates  $c_3$ ,  $s_1$  can be removed
  - $-c_3$ : sum(S.price) ≥ v
- Succinct: Enforce constraint c by explicitly manipulating data
  - S  $\supseteq$  {i-phone, MacAir}
- Convertible: Projection based on the sorted value not sequence order
  - value\_avg(S) < 25; profit\_sum(S) > 160
  - $\max(S)/avg(S) < 2; median(S) \min(S) > 5$

### Timing-Based Constraints in Seq.-Pattern Mining

- Order constraint: Some items must happen before the other
  - {algebra, geometry} → {calculus} (where "→" indicates ordering)
  - Anti-monotonic: Constraint-violating sub-patterns pruned
- Min-gap/max-gap constraint: Confines two elements in a pattern
  - E.g., mingap = 1, maxgap = 4
  - Succinct: Enforced directly during pattern growth
- Max-span constraint: Maximum allowed time difference between the 1<sup>st</sup> and the last elements in the pattern
  - E.g., maxspan (S) = 60 (days)
  - Succinct: Enforced directly when the 1<sup>st</sup> element is determined
- Window size constraint: Events in an element do not have to occur at the same time: Enforce max allowed time difference
  - E.g., window-size = 2: Various ways to merge events into elements

#### Episodes and Episode Pattern Mining

- Episodes and regular expressions: Alternative to seq. patterns
  - Serial episodes: A → B
  - Parallel episodes: A | B | Indicating partial order relationships
  - Regular expressions: (A|B)C\*(D  $\rightarrow$  E)
- Methods for episode pattern mining
  - Variations of Apriori/GSP-like algorithms
  - Projection-based pattern growth
    - Q₁: Can you work out the details?
  - $\Omega_2$ : What are the differences between mining episodes and constraint-based pattern mining?

#### Advanced Frequent Pattern Mining

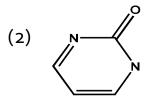
- Mining Diverse Patterns
- Constraint-Based Frequent Pattern Mining
- Mining High-Dimensional Data and Colossal Patterns
- Sequential Pattern Mining
- Graph Pattern Mining

### Frequent (Sub) Graph Patterns

- Given a labeled graph dataset D = {G<sub>1</sub>, G<sub>2</sub>, ..., G<sub>n</sub>), the supporting graph set of a subgraph g is D<sub>q</sub> = {G<sub>i</sub> |  $g \subseteq G_i$ , G<sub>i</sub>  $\in$  D}.
  - support(g) =  $|D_q|/|D|$
- A (sub)graph g is **frequent** if  $support(g) \ge min\_sup$  Ex.: Chemical structures
- Alternative:
  - Mining frequent subgraph patterns from a single large graph or network

 $min_sup = 2$ 

#### Frequent Graph Patterns



#### Applications of Graph Pattern Mining

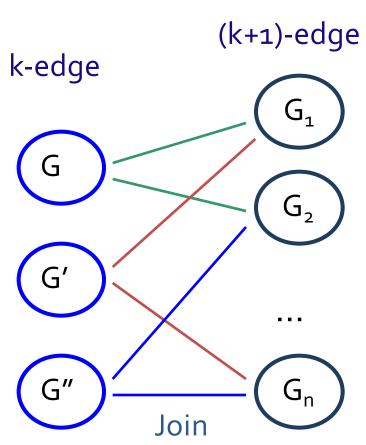
- Bioinformatics
  - Gene networks, protein interactions, metabolic pathways
- Chem-informatics: Mining chemical compound structures
- Social networks, web communities, tweets, ...
- Cell phone networks, computer networks, ...
- Web graphs, XML structures, semantic Web, information networks
- Software engineering: program execution flow analysis
- Building blocks for graph classification, clustering, compression, comparison, and correlation analysis
- Graph indexing and graph similarity search

# Graph Pattern Mining Algorithms: Different Methodologies

- Generation of candidate subgraphs
  - Apriori vs. pattern growth (e.g., FSG vs. gSpan)
- Search order
  - Breadth vs. depth
- Elimination of duplicate subgraphs
  - Passive vs. active (e.g., gSpan (Yan&Han'o2))
- Support calculation
  - Store embeddings (e.g., GASTON (Nijssen&Kok'o4, FFSM (Huan, et al.'o3), MoFa (Borgelt and Berthold ICDM'o2))
- Order of pattern discovery
  - Path → tree → graph (e.g., GASTON (Nijssen&Kok'o4)

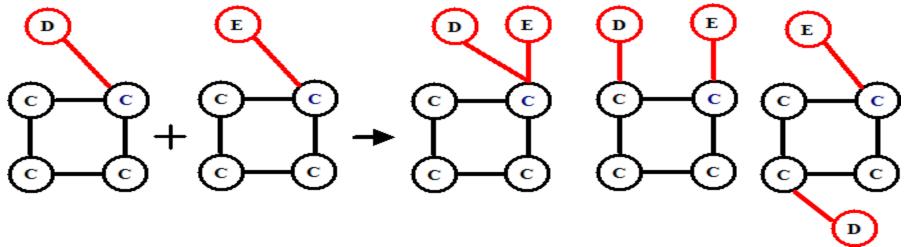
### Apriori-Based Approach

- The Apriori property (antimonotonicity): A size-k subgraph is frequent if and only if all of its subgraphs are frequent
- A candidate size-(k+1) edge/vertex subgraph is generated if its corresponding two k-edge/vertex subgraphs are frequent
- Iterative mining process:
  - Candidate-generation →
     candidate pruning → support
     counting → candidate elimination



### Candidate Generation: Vertex Growing vs. Edge Growing

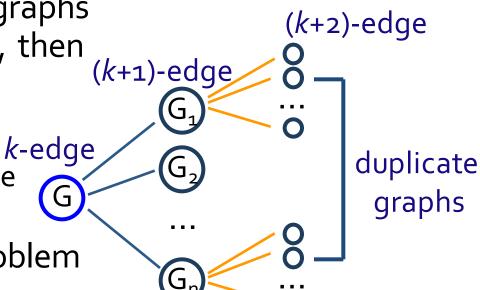
- Methodology: breadth-search, Apriori joining two size-k graphs
  - Many possibilities at generating size-(k+1) candidate graphs



- Generating new graphs with one more vertex
  - AGM (Inokuchi, et al., PKDD'oo)
- Generating new graphs with one more edge
  - FSG (Kuramochi and Karypis, ICDM'01)
- Performance shows via edge growing is more efficient

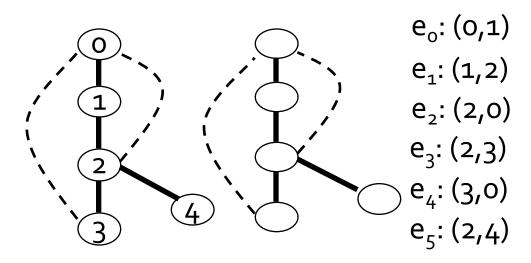
### Pattern-Growth Approach

- Depth-first growth of subgraphs from k-edge to (k+1)-edge, then (k+2)-edge subgraphs
- Major challenge
  - Generating many duplicate subgraphs
- Major idea to solve the problem
  - Define an order to generate subgraphs
  - DFS spanning tree: Flatten a graph into a sequence using depth-first search
  - gSpan (Yan & Han: ICDM'02)



### gSPAN: Graph Pattern Growth in Order

- Right-most path extension in subgraph pattern growth
  - Right-most path: The path from root to the right-most leaf (choose the vertex w. the smallest index at each step)
  - Reduce generation of duplicate subgraphs
- Completeness: The Enumeration of graphs using right-most path extension is <u>complete</u>
- DFS Code: Flatten a graph into a sequence using depth-first search



### Why Mining Closed Graph Patterns?

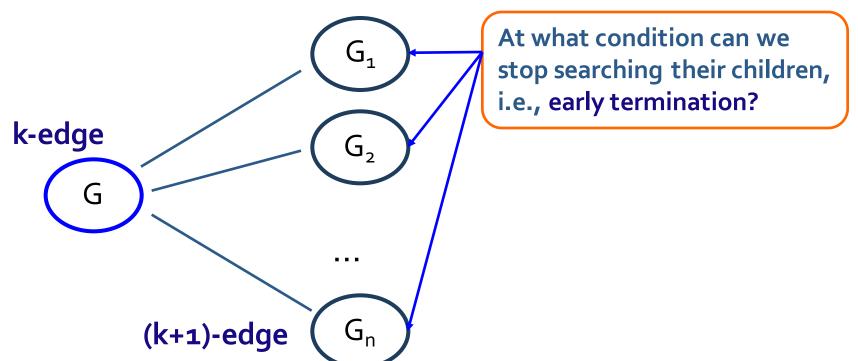
- Challenge: An **n**-edge frequent graph may have 2<sup>n</sup> subgraphs
- Motivation: Explore *closed frequent subgraphs* to handle graph pattern explosion problem
- A frequent graph G is *closed* if there exists no supergraph of G that carries the same support as G

If this subgraph is *closed* in the graph dataset, it implies that none of its frequent super-graphs carries the same support

- Lossless compression: Does not contain non-closed graphs, but still ensures that the mining result is complete
- Algorithm CloseGraph: Mines closed graph patterns directly

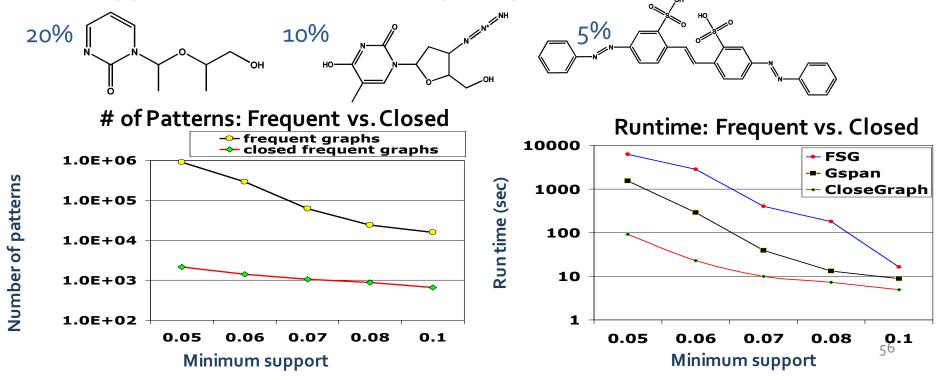
### CloseGraph: Directly Mining Closed Graph Patterns

- CloseGraph: Mining closed graph patterns by extending gSpan
- Suppose G and G<sub>1</sub> are frequent, and G is a subgraph of G<sub>1</sub>
- If in any part of the graph in the dataset where G occurs, G<sub>1</sub> also occurs, then we need not grow G (except some special, subtle cases), since none of G's children will be closed except those of G<sub>1</sub>



# Experiment and Performance Comparison

- The AIDS antiviral screen compound dataset from NCI/NIH
- The dataset contains 43,905 chemical compounds
- Discovered Patterns: The smaller minimum support, the bigger and more interesting subgraph patterns discovered



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