

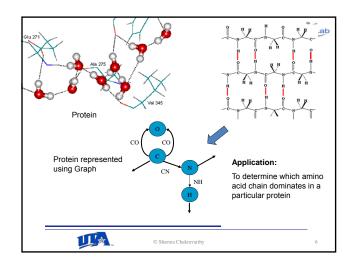
## Need for Graph Mining



- > Significant work in this area includes
  - Subdue substructure discovery algorithm (Cook & Holder),
  - HDB-Subdue (Chakrvarthy, Padmanabhan),
  - Apriori graph mining (AGM) (Inokuchi, Washio, and Motoda),
  - the frequent subgraph (FSG) technique (Karypis & Kuramochi), and
  - gSpan approach (J. Han), also SPIN (Huan, Wang, Prins, and Yang)
- > PageRank and HITS are also graph based



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## **Application Domains**



- > Chemical Reaction chains
- CAD Circuit Analysis
- Social Networks
- Credit Domains
- Web analysis
- Games (Chess, Tic Tac toe)
- Program Source Code analysis
- ➤ Chinese Character data bases
- Geology
- Web and social network analysis



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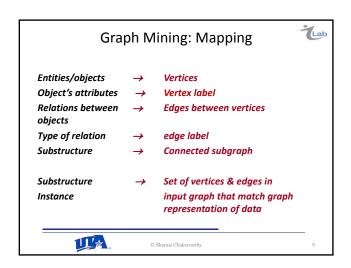
## Graph Based Data Mining

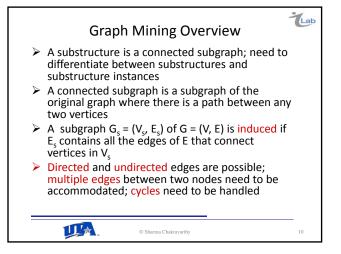


- A Graph representation is an intuitive and an obvious choice for a database that has structural information
- Graphs can be used to accurately model and represent scientific data sets. Graphs are suitable for capturing arbitrary relations between the various objects.
- Graph based data mining aims at discovering interesting and repetitive patterns within these structural representations of data.

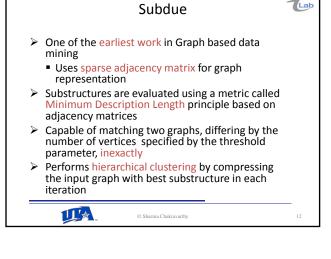


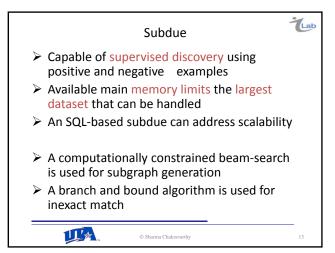
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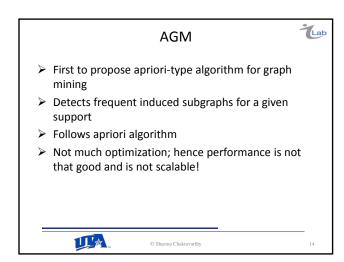


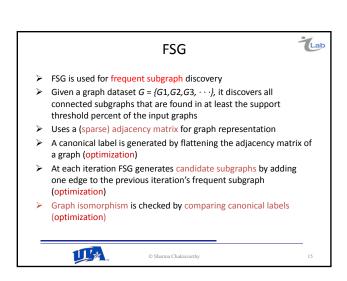


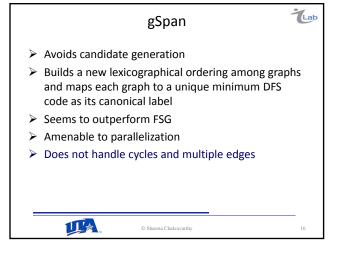
## **Graph Mining: Complexity** > Enumerating all the substructures of a graph has exponential complexity Subgraph isomorphism (or subgraph matching) is NP-complete However, graph isomorphism although belongs to NP is neither known to be solvable in polynomial time nor NP-complete ➤ Generating canonical labels is O(|V|!), where V is the number of vertices All approaches have to deal with the above in order to be able to work on large data sets Different approaches do it differently; scalability depends on the approach and the use of representation © Sharma Chakravarthy

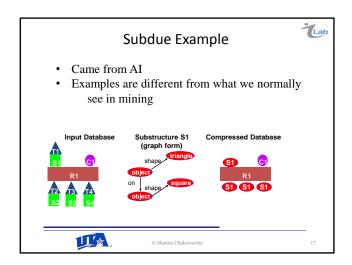


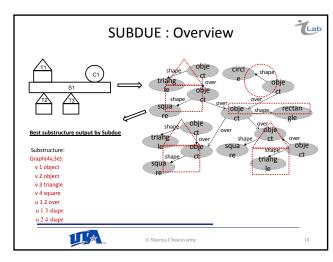


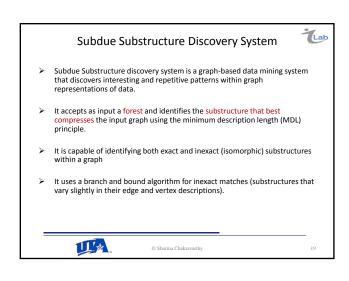


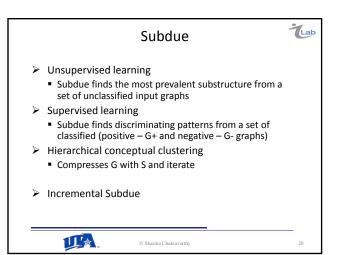


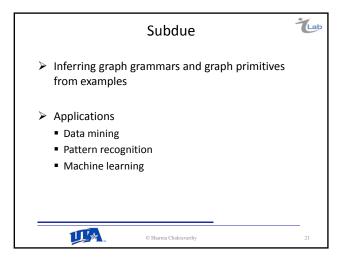


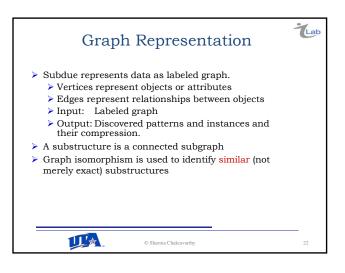


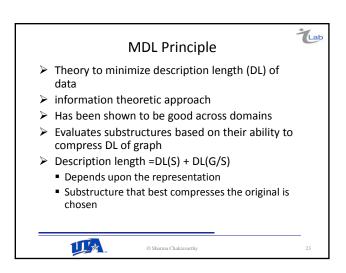


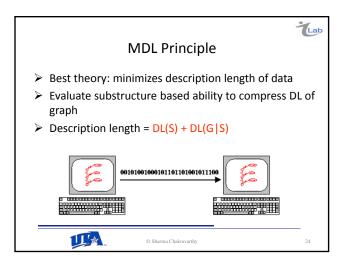


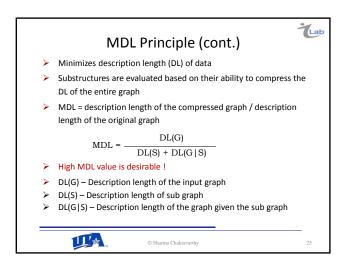


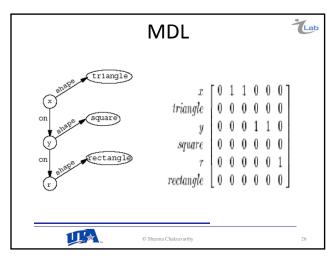


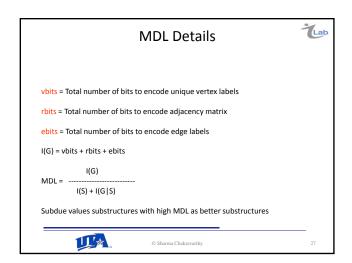


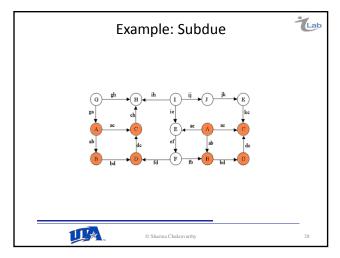


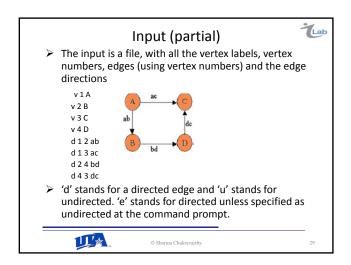


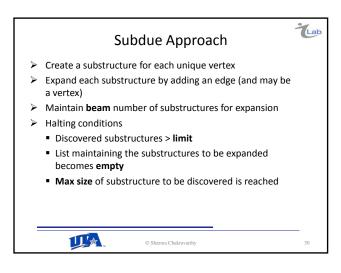


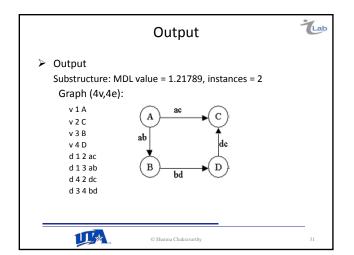


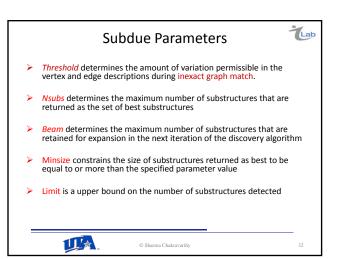


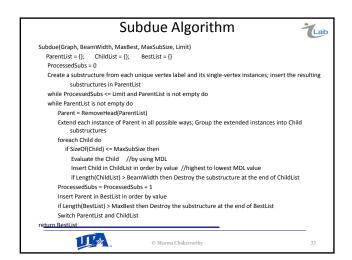


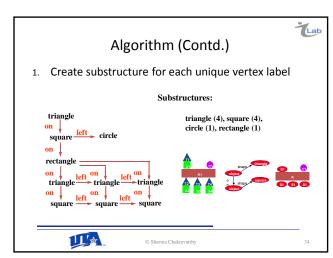


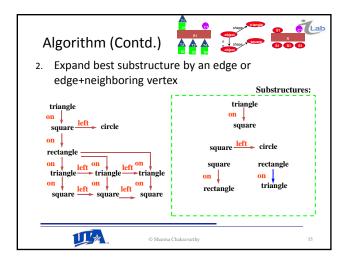


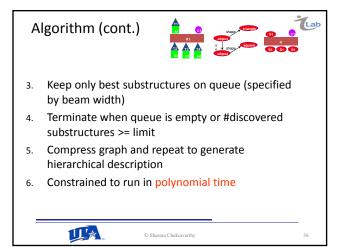


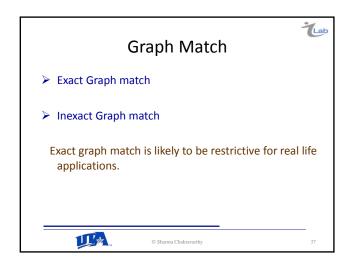


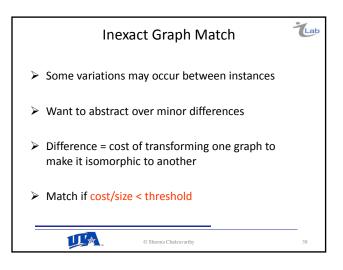


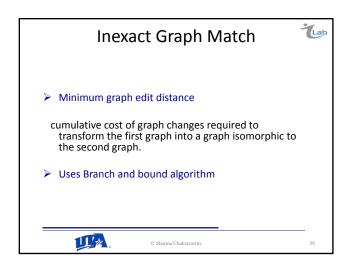


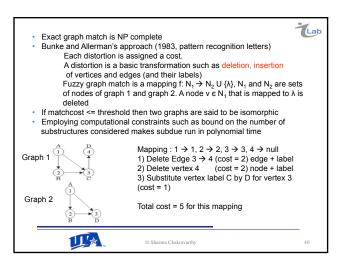


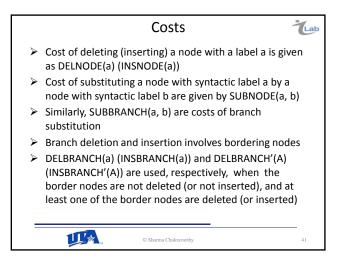


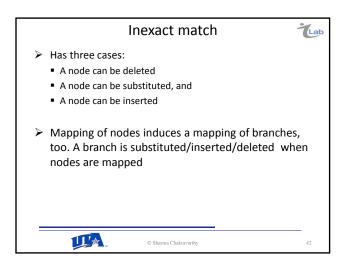


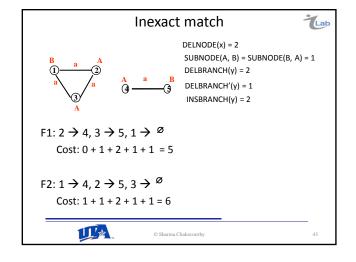


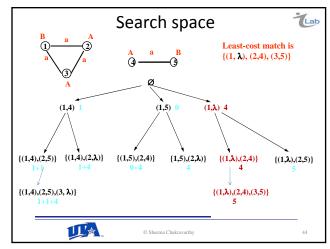


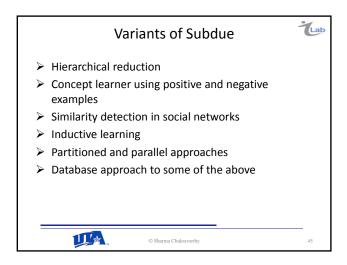


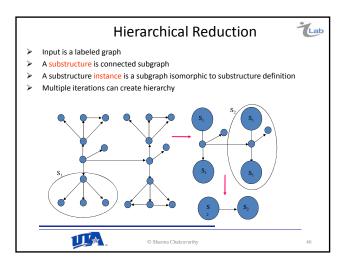


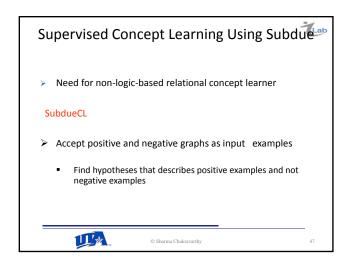


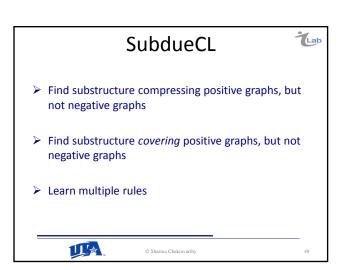


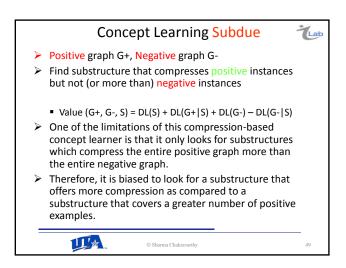


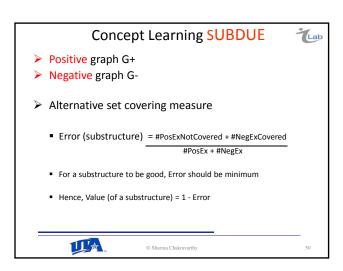




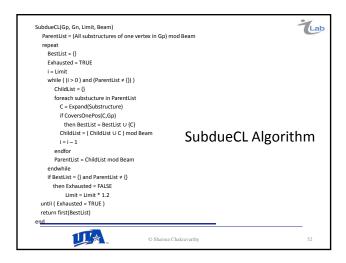


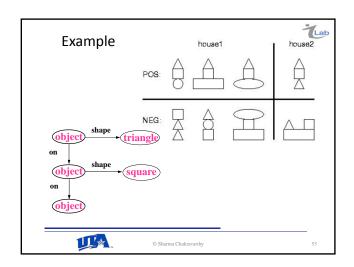


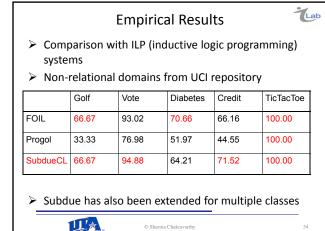


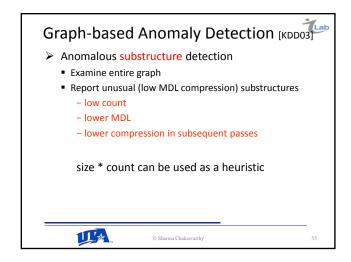


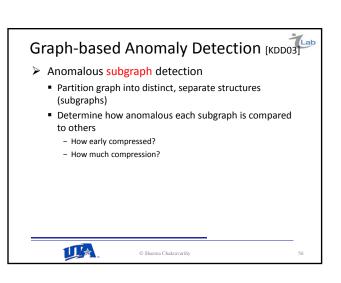
```
Hypotheses detection using coverage *Lab
Main(Gp, Gn, Beam, Limit)
  H = \{\};
  repeat
   repeat
      BestSub = SubdueCL(Gp, Gn, Beam, Limit)
       if BestSub = {}
           then Beam m= Beam * 1.1
   until (BestSub <> {})
   Gp = Gp - {p in Gp | BestSub covers p}
   H = H + BestSub
  until Gp = {}
  return H
end
         111/2
                          © Sharma Chakravarthy
```

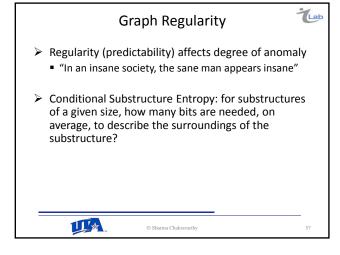


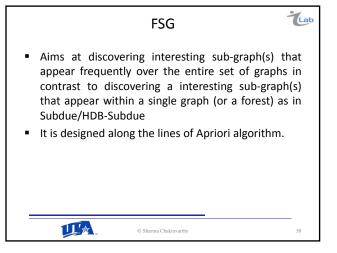


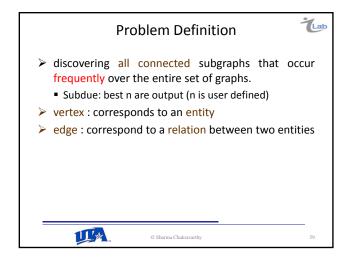


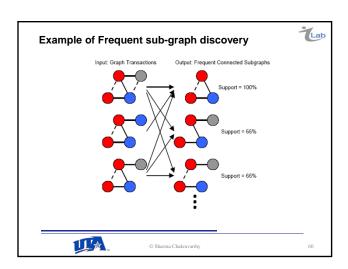


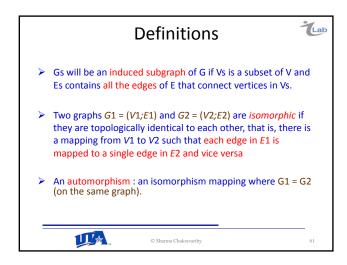


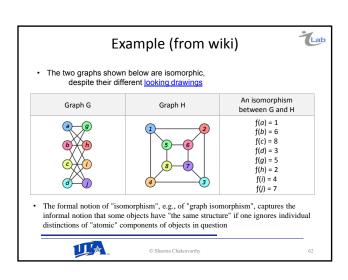


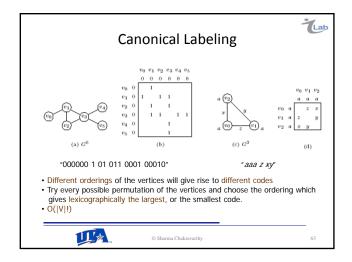


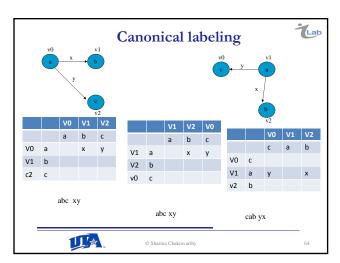


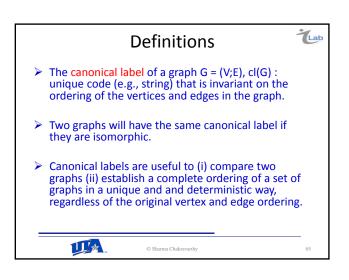


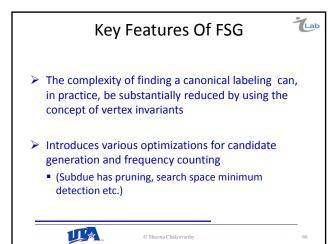


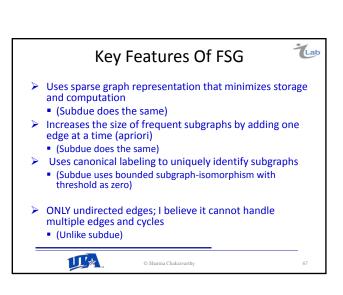


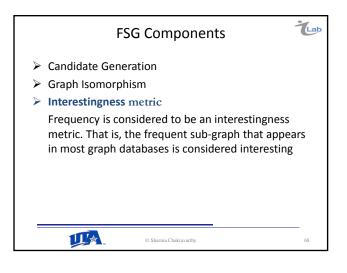


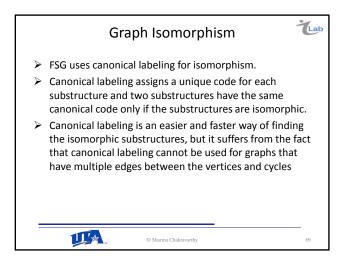


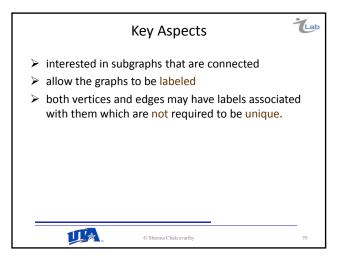


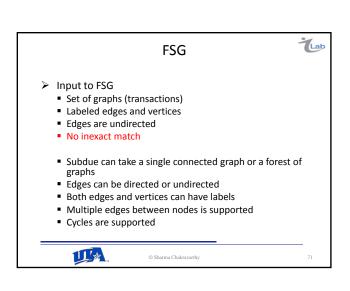


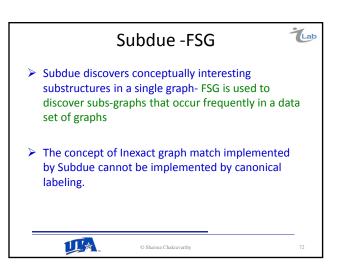
















- Subdue requires to know the graph structure in the candidate generation phase but FSG does not need the adjacency list in the candidate generation phase, candidates are generated using the existing frequent substructures.
- FSG prunes the candidate substructures using the 'downward closure property', which allows a k+1 size substructure to be a candidate only if all its k subs-graphs are frequent substructures.
- Grows Vs. generates



© Character

## Subdue - FSG



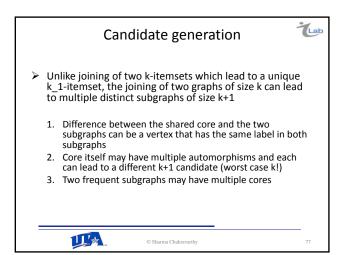
- > Frequent subgraphs are found based on the set covering approach (frequency)
  - In Subdue subgraphs are found based on MDL (the graph that minimizes the description length of the input)
- User defined support threshold minimum percentage of graphs in which a subgraph has to be found

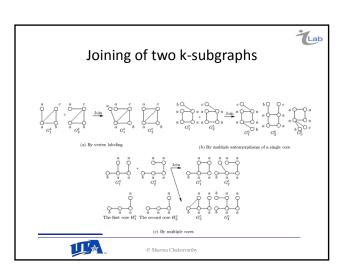


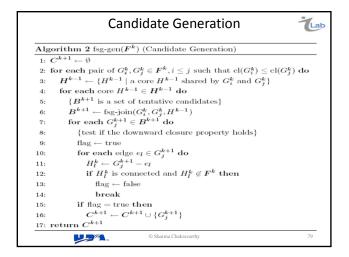
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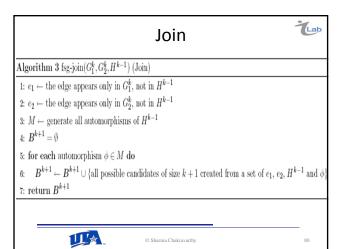
### Algorithm fsg(D; t) 1: F(1) = detect all frequent 1-subgraphs in D 2: F(2) = detect all frequent 2-subgraphs in D 3: k = 3 4: while F (k-1) != NULL; do 5: C(k) = fsg-gen(F(k-1)) 6: for each candidate G(k) in C(k) do G(k).count = 0 for each transaction T in D do 8: if candidate G(k) is included in transaction T then 9. 10: G(k).count++ 11: $F(k) = \{G(k) \text{ in } C(k) \mid G(k).count >= t \mid D \mid \}$ 12: k++ 13: return F(1);F(2); ...... ;F(k-2) © Sharma Chakravarthy

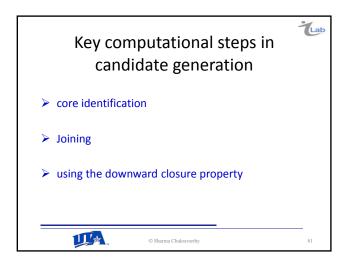
# Candidate generation Candidates are the substructures which would be searched and counted in the given graph databases create a set of candidates of size k+1, given frequent k-subgraphs. by joining two frequent k-subgraphs (using downward closure property) must contain the same (k-1)-subgraph (common core) Self-join required for unlabeled graphs Subdue extends subgraph in every possible way via an edge and a vertex

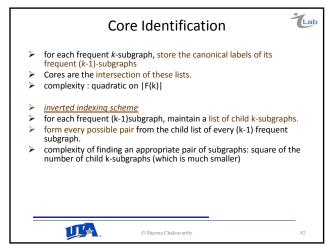


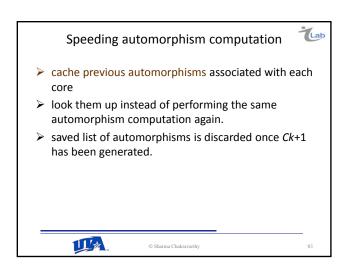


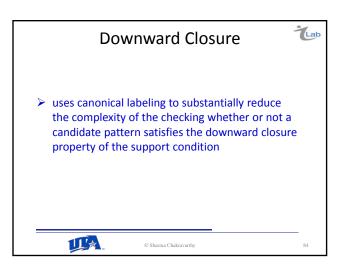


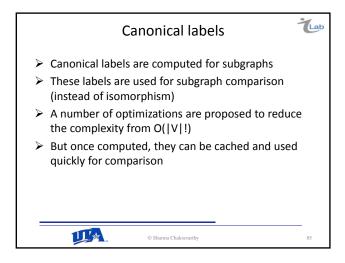


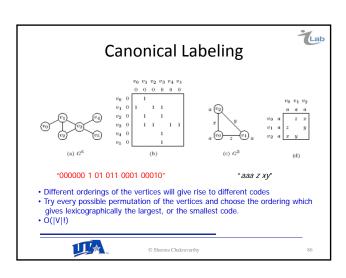


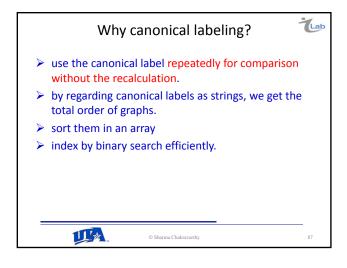


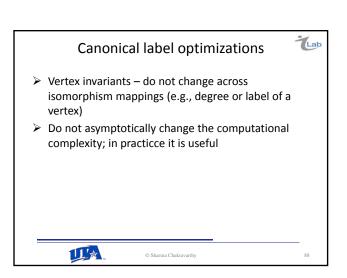


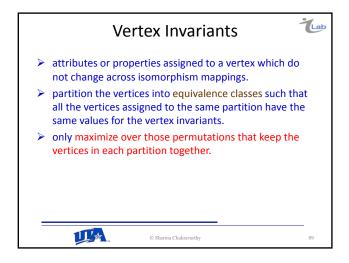


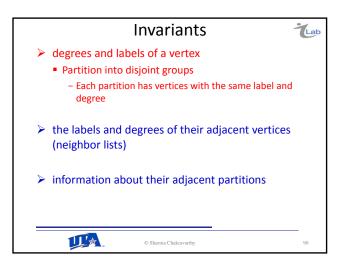


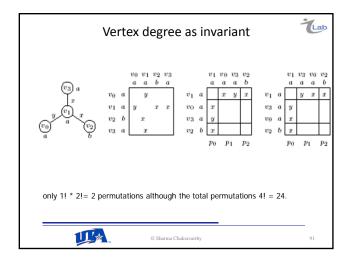


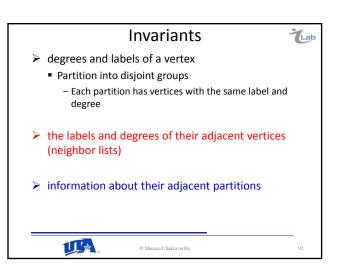


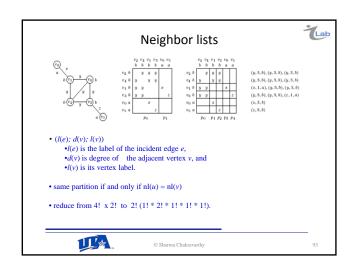


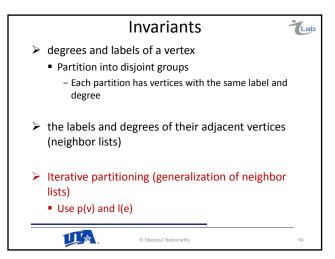


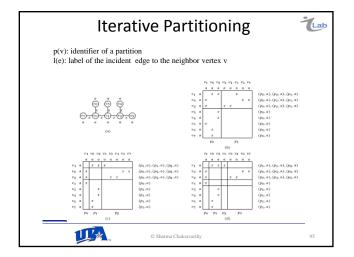


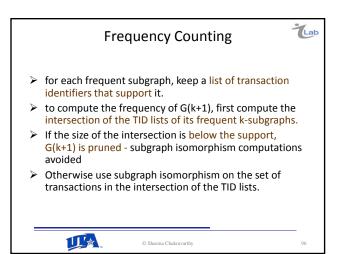


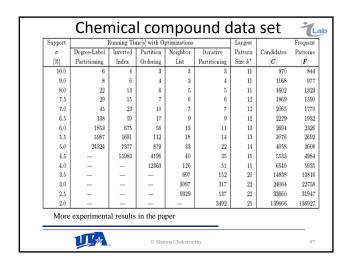


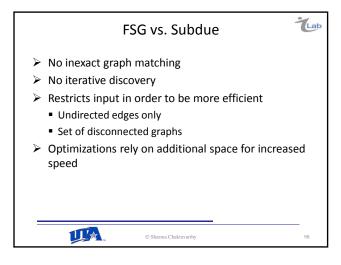


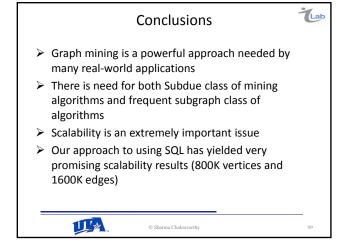




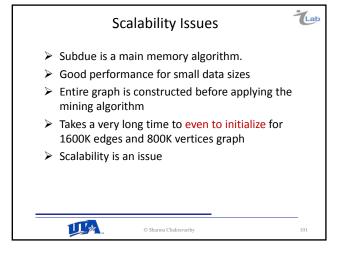


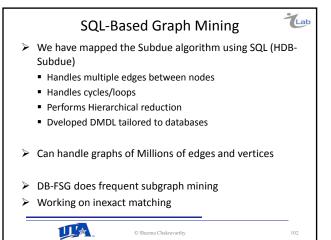


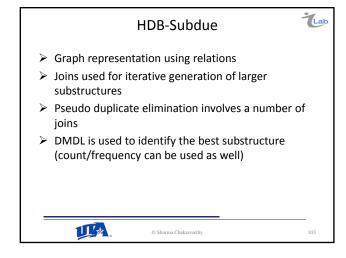


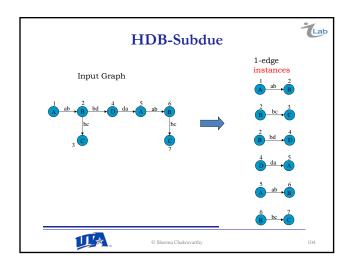


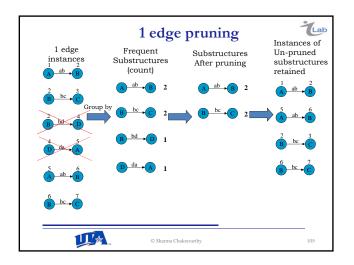
Comparison										
	Subdue	FSG	AGM	gSpan	HDBSubdue					
Graph Mining	<b>✓</b>	<b>✓</b>	✓	<b>✓</b>	✓					
Multiple edges	✓	×	*	×	✓					
Hierarchical reduction	✓	×	*	×	✓					
Cycles	✓	<b>✓</b>	<b>√</b>	×	✓					
Evaluation metric	MDL	Frequency	Support, Confidence	Frequency	DMDL (frequency)					
Inexact graph match With threshold	✓	×	×	×	×					
Memory limitation	✓	✓	✓	✓	×					
<u> </u>	100									

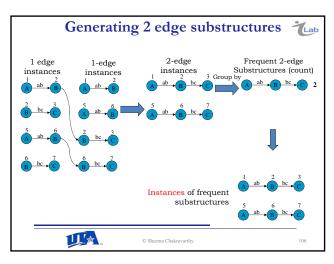


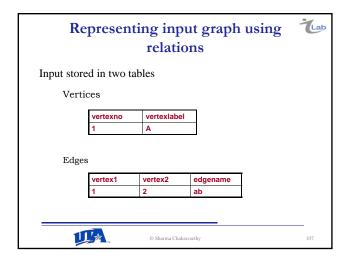


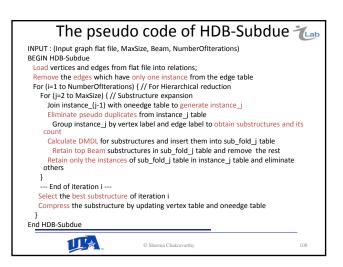


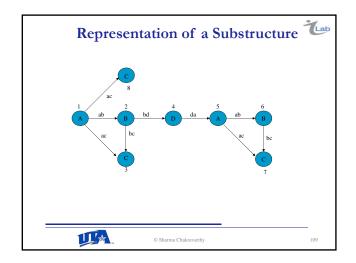


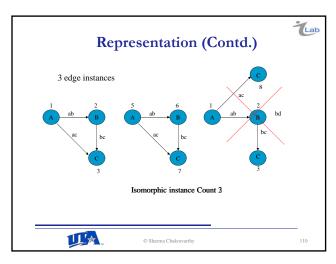


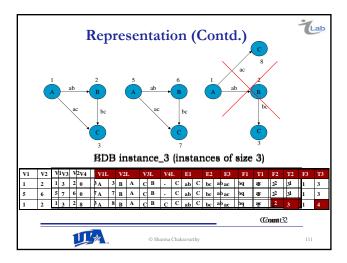


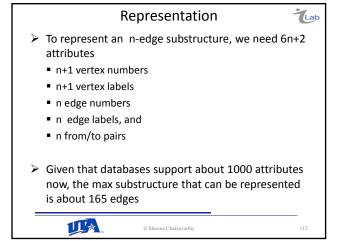


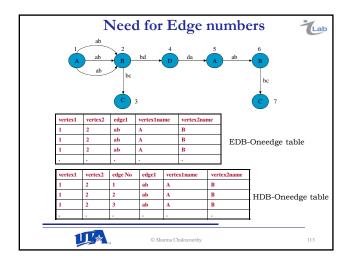


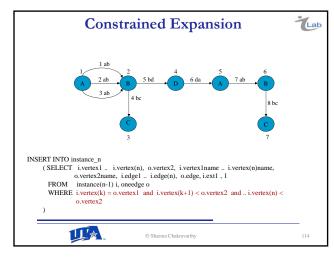


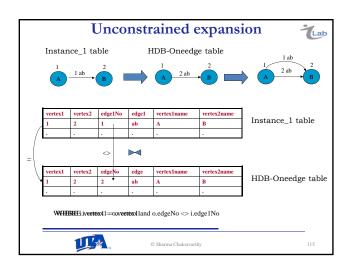


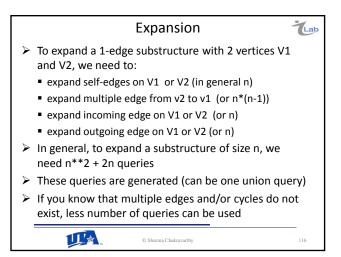




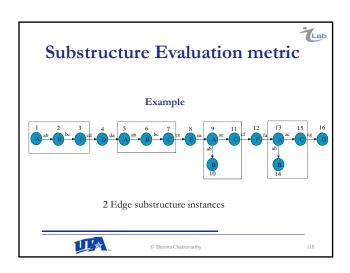


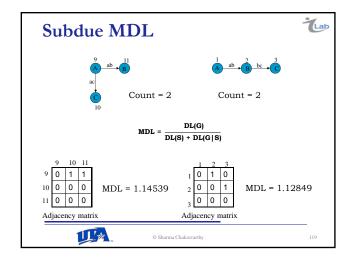


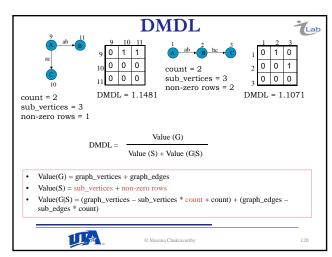


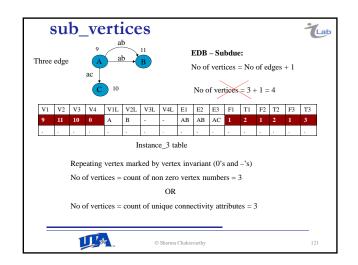


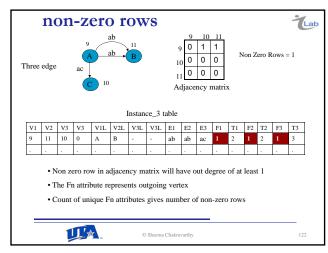
Expansion									Lab	
VL1	VL2	VL3	EL1	EL2	F1	T1	F2	T2 (	COUNT	DMDL
Α	В	С	AB	AC	1	2	1	3	3	1.8
Α	В	D	AB	BD	1	2	2	3	1	0.9
В	D	Α	BD	DA	1	2	2	3	1	0.9
D	Α	В	DA	AB	1	2	2	3	1	0.9
D	Α	С	DA	AC	1	2	2	3	1	0.9
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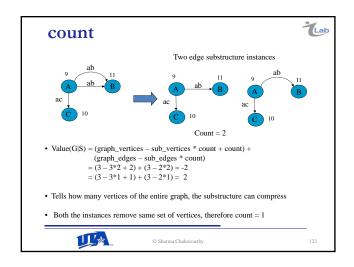


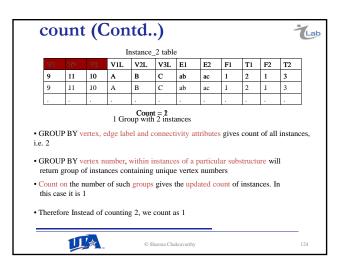


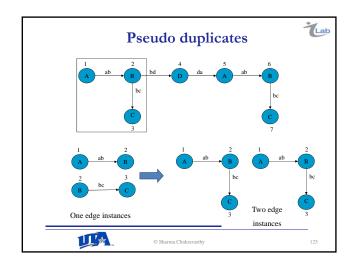


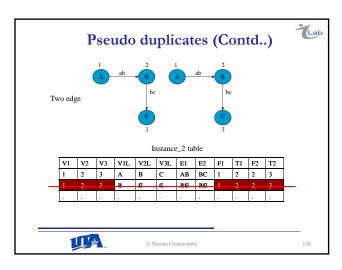


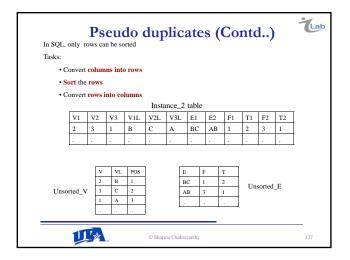


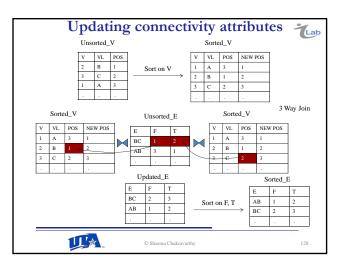


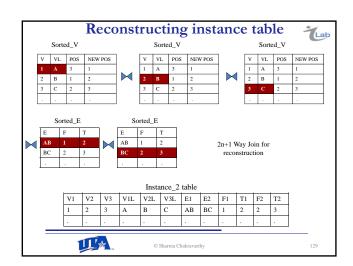


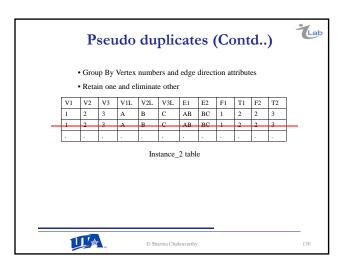


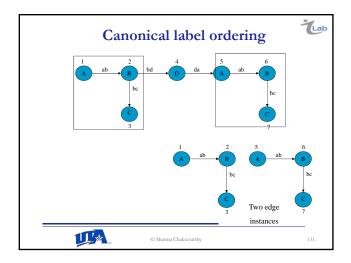


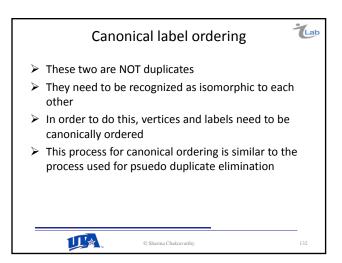


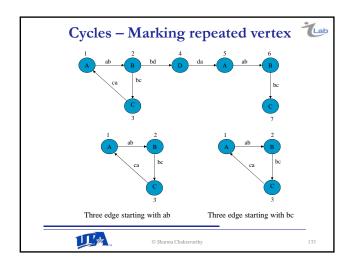


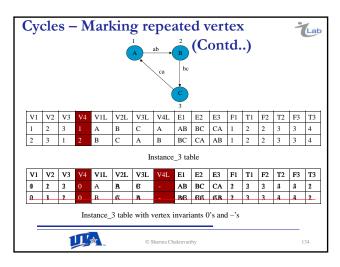


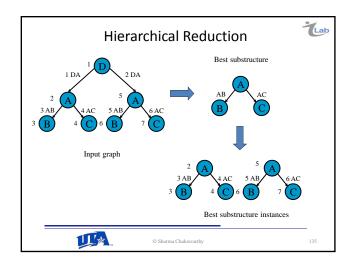


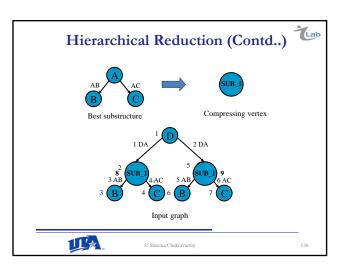


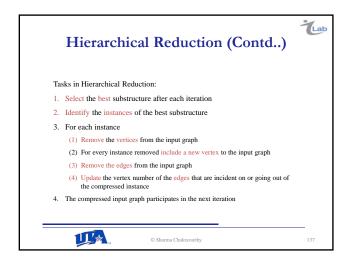


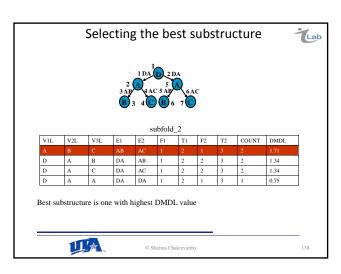


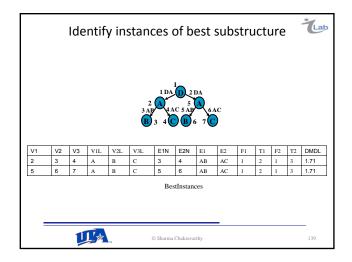


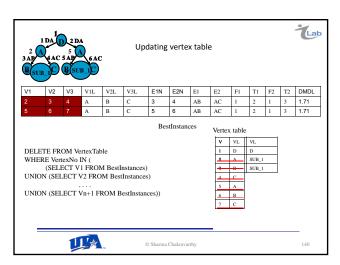


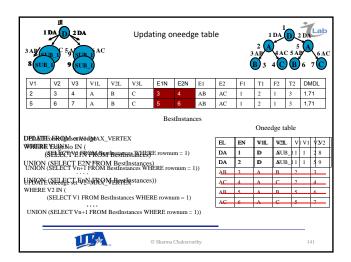


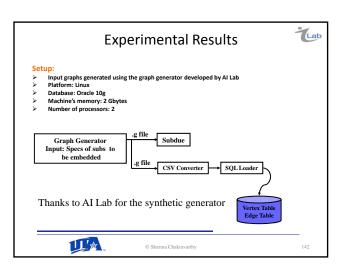


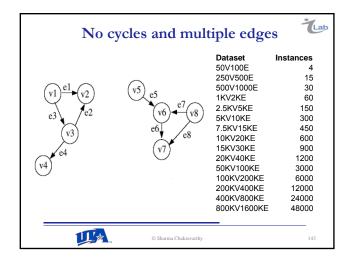


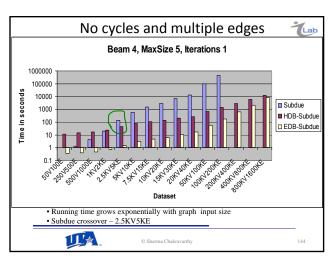


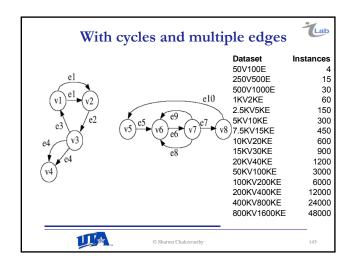


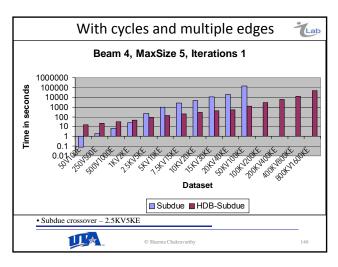


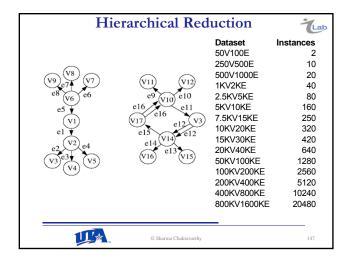


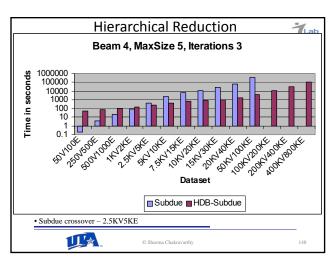


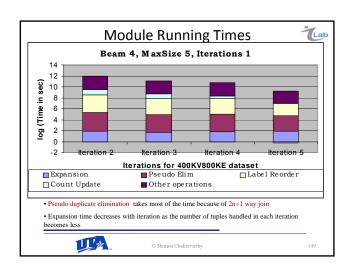


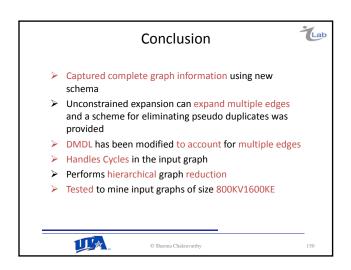


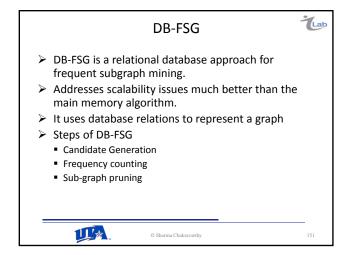


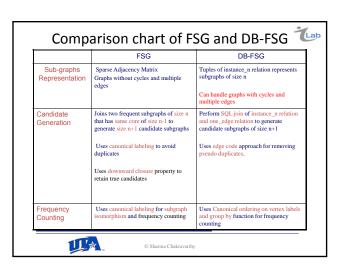


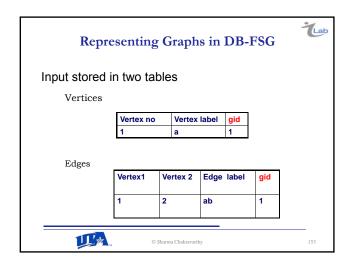


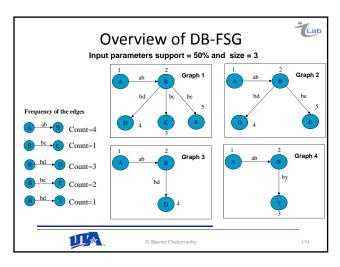


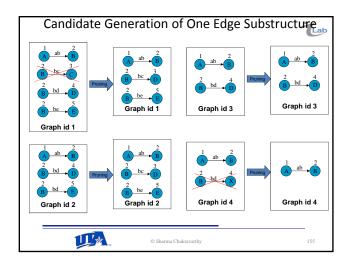


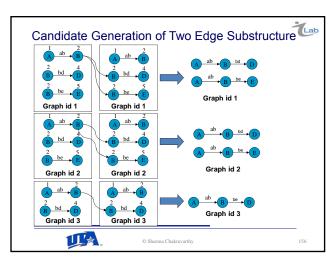


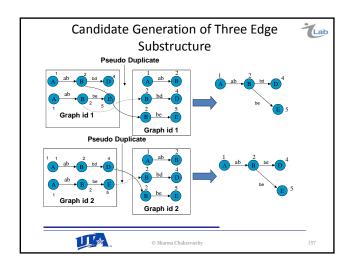


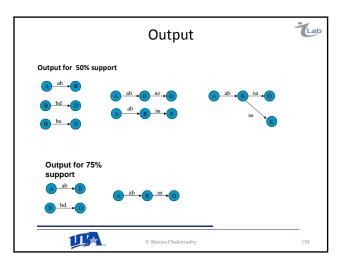


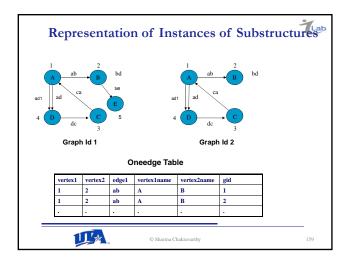


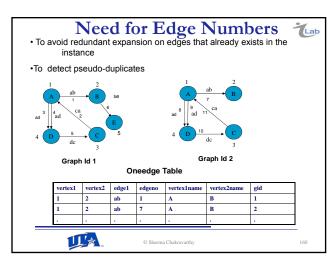


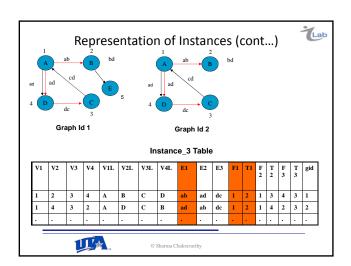


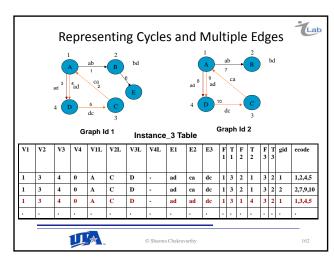












# Detecting Pseudo duplicates using Edge Code

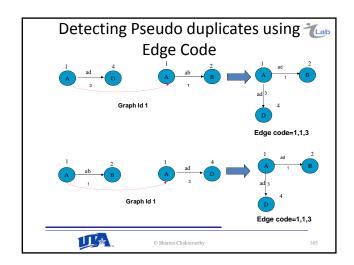
- > Each edge in a graph has unique edge number
- > All pseudo duplicates have same edges and edge number in different order.
- Hence, we can construct a unique code based on edge numbers and gid
- Edge code is a string formed by concatenating gid with edge numbers sorted in ascending order and separated by comma

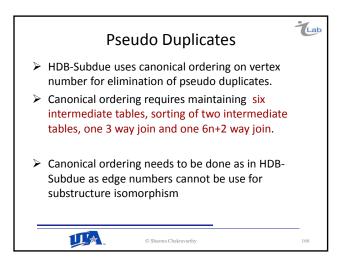


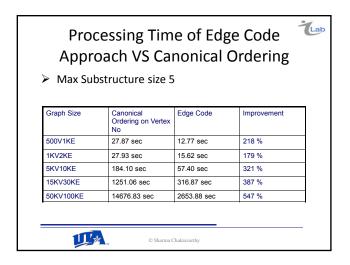
# Detecting Pseudo duplicates using Edge Code

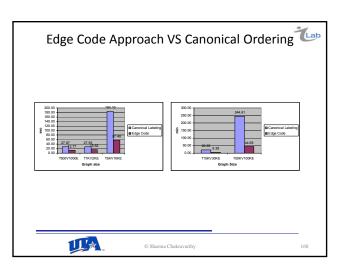
- ➤ If we have edge code for instance of size n then constructing edge code for instance of size n+1 expanded from the instance is just placing the new edge number in the proper position in edge code
- Hence the complexity is O(n) for constructing edge code for n+1 size instance from expanded from n sized instance



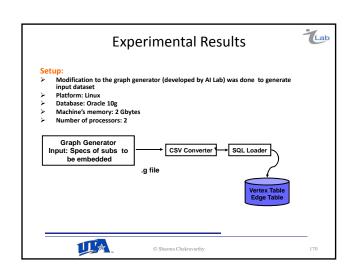


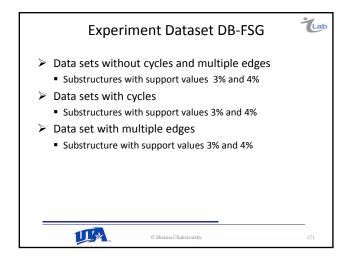


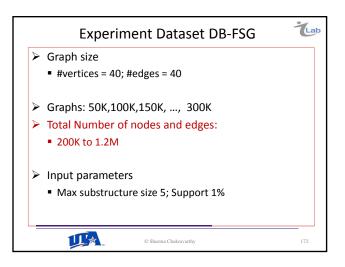


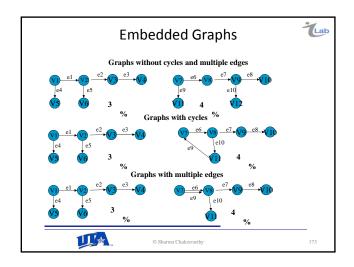


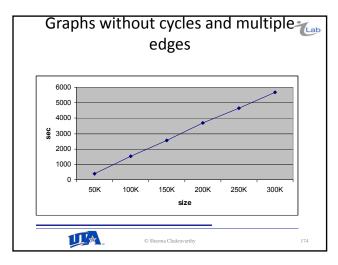
# Frequency Counting and Substructure Pruning Support count=(support X #Graphs)/100 For each graph only one instance per substructure is included for frequency counting of substructures Instances of substructure with frequency more than or equal to support count is retained.

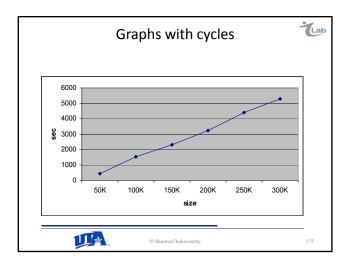


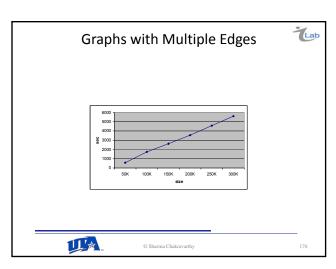


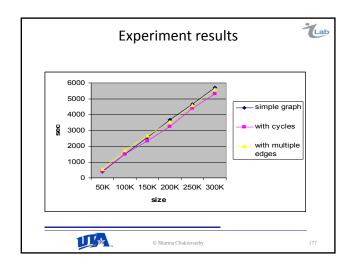


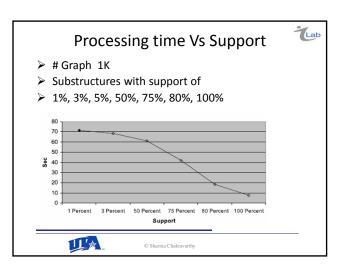


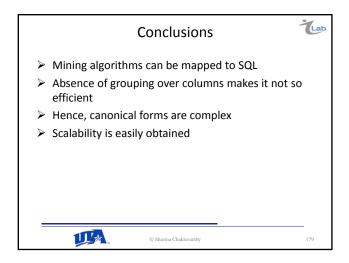


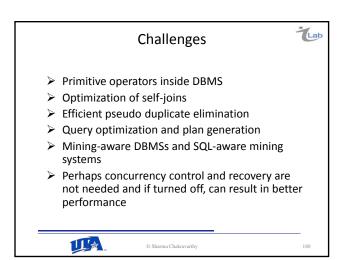












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