

Minimum spanning trees

Data Structures and Algorithms for Computational Linguistics III
(ISCL-BA-07)

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So far...

a quick summary

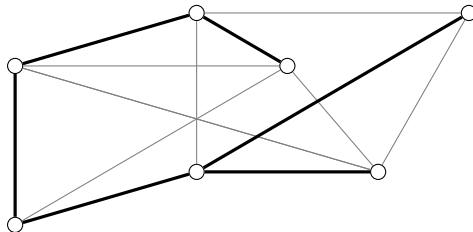
- Recap: arrays, lists, queues, stacks, ...
- Common algorithmic patterns: recursion, brute force, divide and conquer, dynamic programming, greedy algorithms
- Analysis of algorithms: asymptotic, average/worst case analysis, big-O, big- Ω , big- Θ , complexity classes
- Sorting: insertion sort, quicksort, merge sort
- Trees: ordered trees, binary trees, tree traversals, ...
- Priority queues and heaps, heap sort
- Graphs: graph traversal, directed graphs

Spanning trees

reminder

A spanning tree of a graph is

- A *spanning subgraph*: it includes all nodes
- It is a *tree*: it is *acyclic*, and *connected*

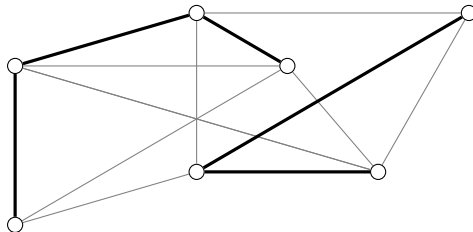


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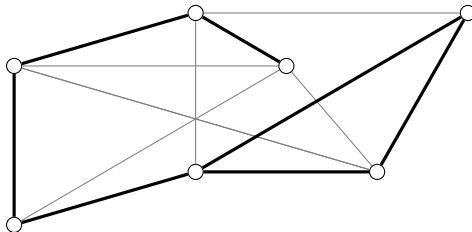


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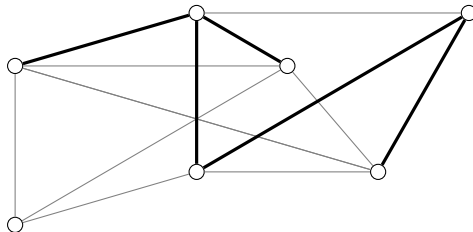


Spanning trees

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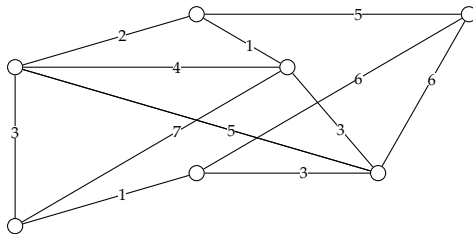
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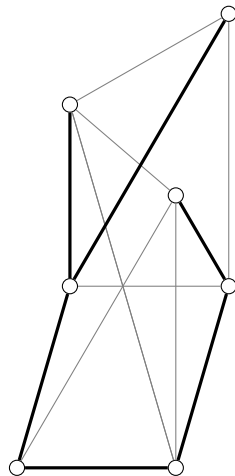
Weighted graphs

- A *weighted graph* is a graph, where each edge is associated with a weight
- Weights can be any numeric value, but for some algorithms require
 - Non-negative weights
 - ‘Euclidean’ weights: weights that are proper distance metrics
- Weights often indicate distance or cost, but they can also represent positive relations (e.g., affinity between nodes)
- Weight of a path is the sum of wights of the edges on the path



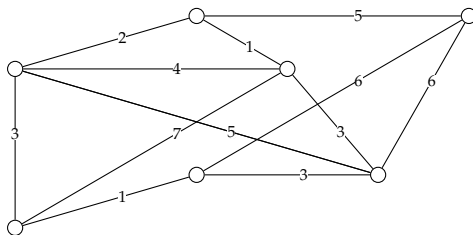
Minimum spanning trees

- A minimum spanning tree (MST) is a spanning tree of weighted graph with minimum total weight
- MST is a fundamental problem with many applications, including
 - Network design (communication, transportation, electrical, ...)
 - Cluster analysis
 - Approximate solutions to traveling salesman problem
 - Object/network recognition in images
 - Avoiding cycles in broadcasting in communication networks
 - Dithering in images, audio, video
 - Error correction codes
 - DNA sequencing
 - ...



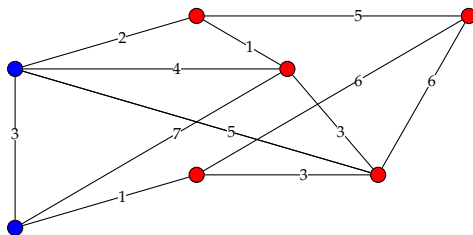
The 'cut property'

- A *cut* of a graph is a partition that divides its nodes into two disjoint (non-empty) sets
- Given any cut, the edge with the lowest weight across the cut is in the MST



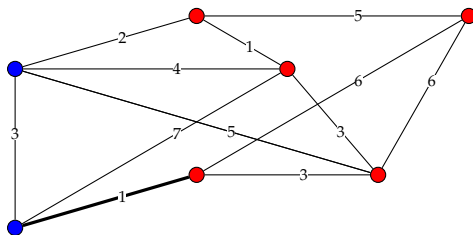
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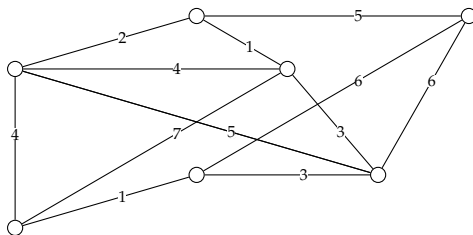
Prim-Jarník algorithm

intuition

- Prim-Jarník algorithm is a greedy algorithm for finding an MST for a weighted undirected graph
- Algorithm starts with a single 'start' node, and grows the MST greedily
- At each step we consider a cut between nodes visited and the rest of the nodes, and select the minimum edge across the cut
- Repeat the process until all nodes are visited

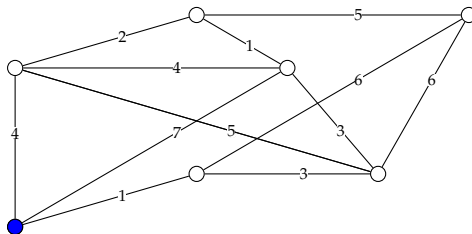
Prim-Jarník algorithm

demonstration



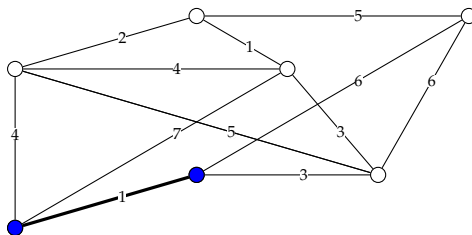
Prim-Jarník algorithm

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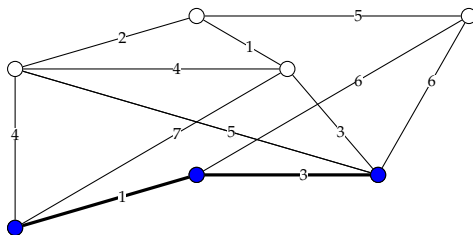
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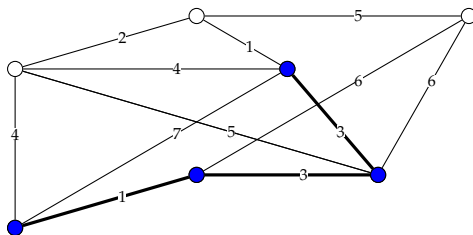
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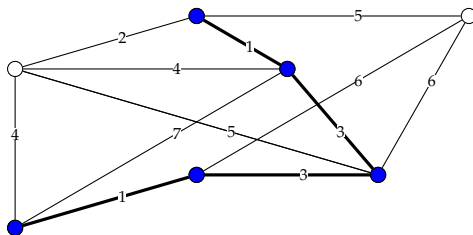
Prim-Jarník algorithm

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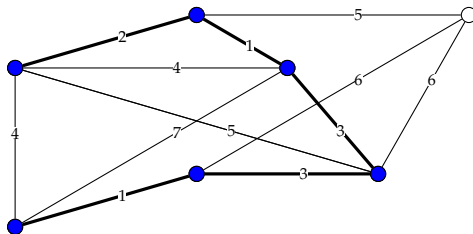
Prim-Jarník algorithm

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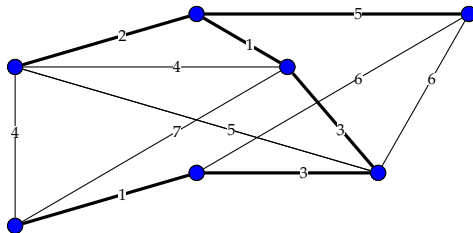
Prim-Jarník algorithm

demonstration



Prim-Jarník algorithm

demonstration



Prim-Jarník algorithm

analysis

- Two loops over number of nodes n , $O(n^2)$ if we need to search
- If we use a priority queue for Q , then complexity becomes $O(m \log m)$

```

1: pick any node  $s$ 
2:  $C[s] \leftarrow 0$ 
3: for each node  $v \neq s$  do
4:    $C[v] \leftarrow \infty$ 
5:    $E[v] \leftarrow \text{None}$ 
6:  $T \leftarrow \emptyset$ 
7:  $Q \leftarrow \text{nodes}$ 
8: while  $Q$  is not empty do
9:   Find the node  $v$  with  $\min C[n]$ 
10:  Connect  $v$  to  $T$ 
11:  for edge  $(v, w)$  in  $Q$  do
12:    if  $\text{cost}(v, w) < C[w]$  then
13:       $C[w] \leftarrow \text{cost}(v, w)$ 
14:       $E[w] \leftarrow v$ 

```

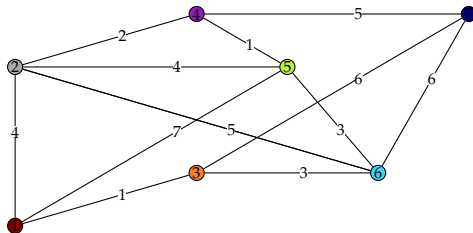
Kruskal's algorithm

intuition

- Another popular algorithm for finding MST on undirected graphs
- The main idea is starting with each node in its own partition
- At each iteration, we choose the edge with the minimum weight across any two clusters, and join them
- Algorithm terminates when there are no clusters to join

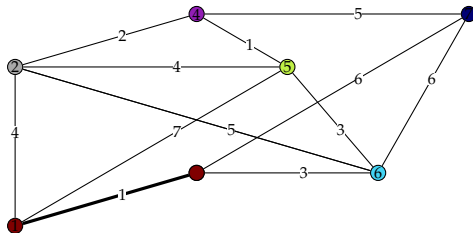
Kruskal's algorithm

demo



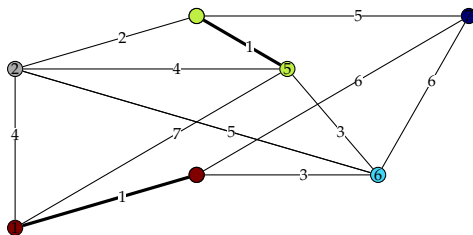
Kruskal's algorithm

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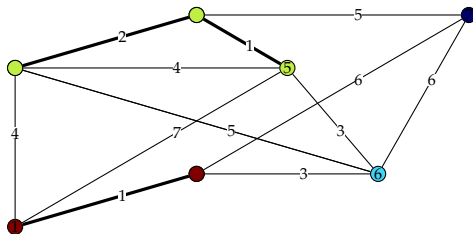
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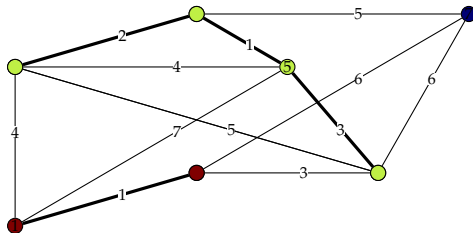
Kruskal's algorithm

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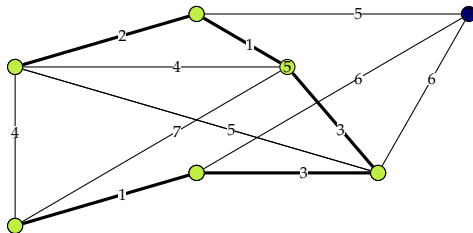
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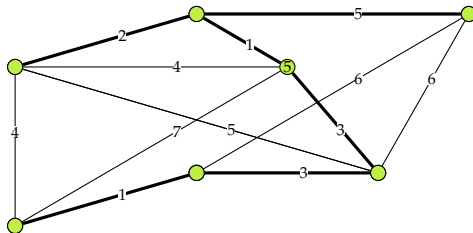
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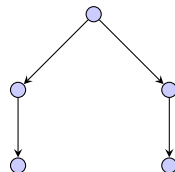
analysis

- Loop over edges, but beware of the sorting requirement
- With simple data structures then complexity is $O(m \log m)$

```
1:  $T \leftarrow \emptyset$ 
2: for each node  $v$  do
3:   create_cluster(v)
4: for  $(u,v)$  in edges sorted by weight do
5:   if  $\text{cluster}(u) \neq \text{cluster}(v)$  then
6:      $T \leftarrow T \cup \{(u, v)\}$ 
7:     union(cluster(u), cluster(v))
```

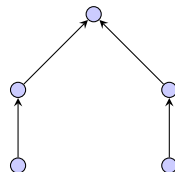
Directed trees

- Trees with directed edges come in few flavors
 - A *rooted directed tree* (arborescence) is an acyclic directed graph where all nodes are reachable from the root node through a single directed path (this is what computational linguists simply calls a *tree*)
 - An anti-arborescence is a rooted directed tree where all edges are reversed
 - A polytree (also called a directed tree) is a directed graph where undirected edges form a tree
- The equivalent of finding an MST in a directed graph is finding a rooted directed tree (arborescence)



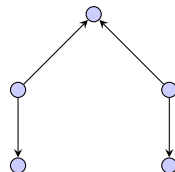
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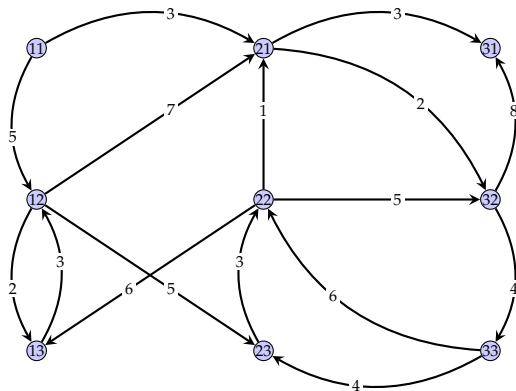
Chu–Liu/Edmonds algorithm

a sketch

- The MST for a directed graph has to start from a designated root node
 - If selected node has any incoming edges, remove them
 - It is also a common practice to introduce an artificial root node with equal-weight edges to all nodes
- For all non-root nodes, select the incoming edge with lowest weight, remove others
- If the resulting graph has no cycles, it is an MST
- If there are cycles break them
 - Consider the cycle as a single node
 - Select the incoming edge that yields the lowest cost if used for breaking the cycle
- Repeat until no cycles remain

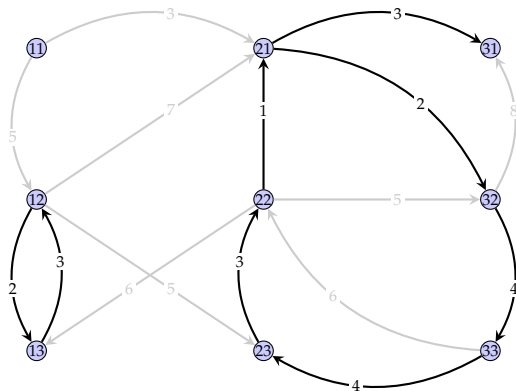
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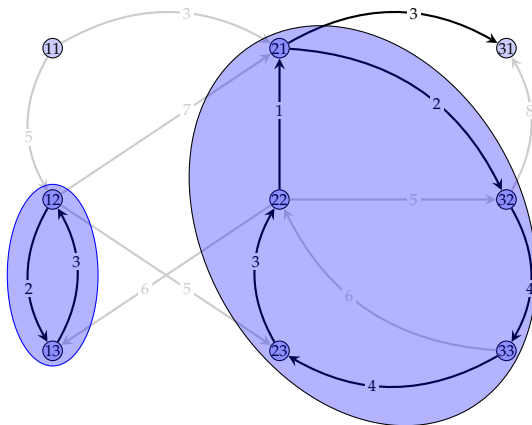
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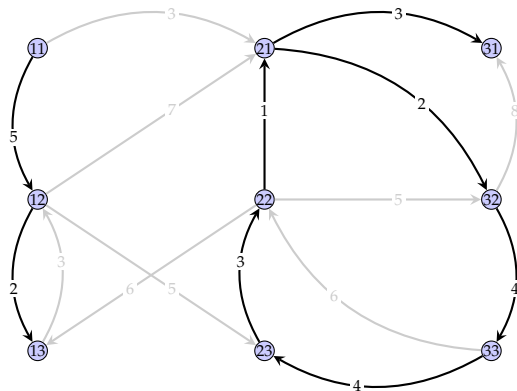
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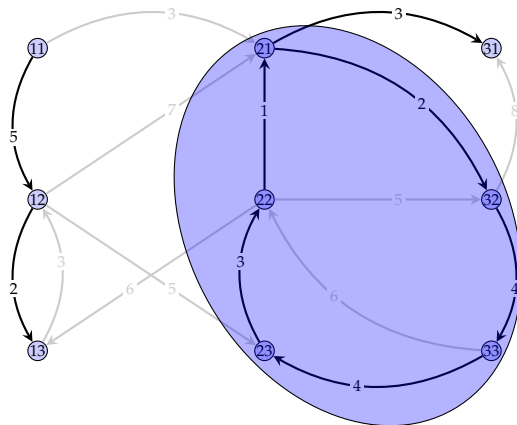
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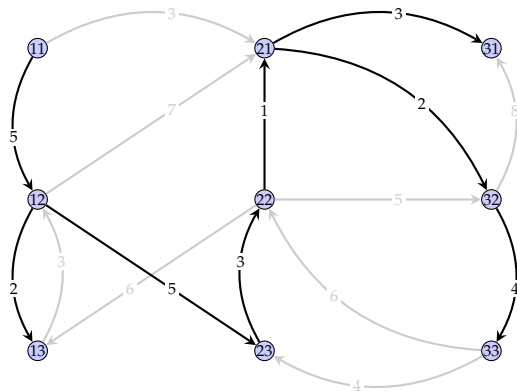
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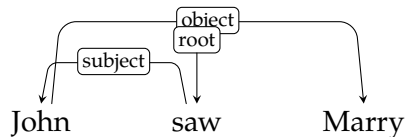
Chu–Liu/Edmonds algorithm

analysis

- The algorithm is generally defined recursively: at each step, create new graph with a contracted cycle call the procedure with the new graph
- At most n recursions: the cycle has to include more nodes at every step
- At each call, m steps for finding minimum incoming edge (also finding a cycle with $O(n)$, but $m \geq n$)
- The ‘vanilla’ algorithm runs in $O(mn)$
- There are improved versions

Chu–Liu/Edmonds algorithm in Computational Linguistics

dependency parsing



- In a dependency analysis, the structure of the sentence is represented by asymmetric binary relations between syntactic units
- Each relation defines one of the words as the head and the other as dependent
- Often an artificial root node is used for computational convenience
- The links (relations) may have labels (dependency types)
- A dependency analysis (parse) is simply a rooted directed tree

Chu–Liu/Edmonds for dependency parsing

- Begin with fully connected weighted graph, except the root node has no incoming edges
- Weights are estimated from a treebank, typically determined by a machine learning method trained on a treebank
- We often use probabilities rather than costs/distances, so, rather than minimizing, maximize the weight of the tree
- Given the fully connected graph, now the parsing becomes finding the MST
- This method is one of the most common (and successful) approaches to dependency parsing

Summary

- Minimum spanning trees have many applications
- An MST of a undirected graph can be found (efficiently) using Prim-Jarník or Kruskal's algorithms
- For directed graph, the corresponding problem can be solved using Chu–Liu/Edmonds algorithm (technically what we find is a rooted directed tree, or arborescence)
- MST also has quite a few applications in CL/NLP

Next:

- Shortest paths

Acknowledgments, credits, references

