Minimum Spanning Trees

COMPSCI 220: WEEK 13.1-13.2

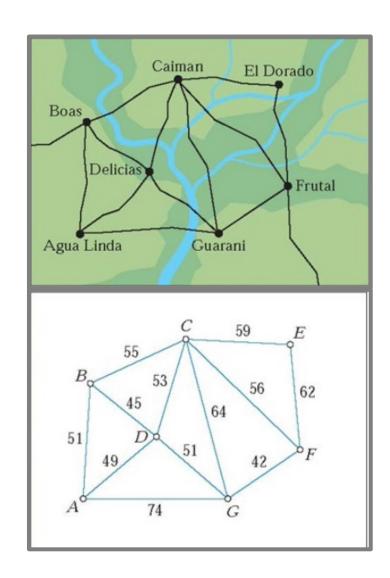
Instructor: Meng-Fen Chiang





OUTLINE

- Terminology
 - Spanning Trees
 - Minimum Spanning Trees
- Finding minimum Spanning Trees Algorithms
 - Prim's Algorithm
 - Kruskal's Algorithm
 - Time Complexity Analysis



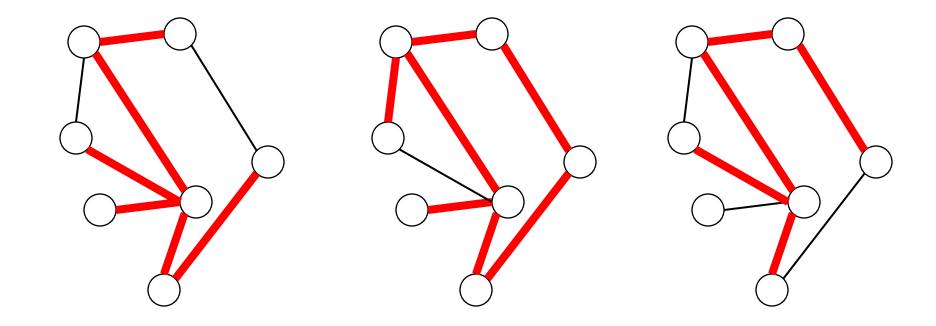


Spanning Trees

- Definition. A spanning tree of a graph G is a connected acyclic graph T such that it contains all the vertices of G, e.g. V(T) = V(G), and a subset of edges
- On a spanning tree T of G, we can walk from any vertex of G to any other vertex of G along exactly one path whose edges are part of the tree T



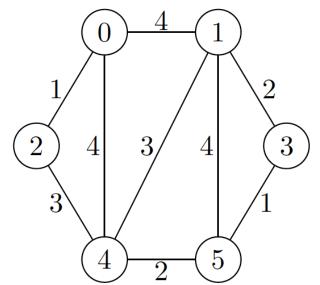
Exercise: Spanning Trees — Yes or No?

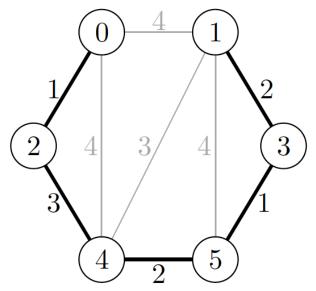




Minimum Spanning Tree Problem

- Minimum spanning tree (MST) problem on a weighted connected graph G:
- Find a spanning tree (subgraph containing all vertices that is a tree) of minimum total weight.



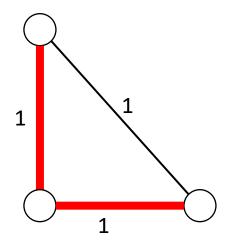


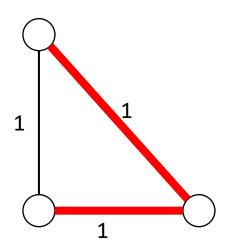
The MST with the total weight 9



Example: Minimum Spanning Trees

• MSTs are not unique







Applications of MSTs

- Design of networks, including computer networks, telecommunications networks, transportation networks, water supply networks, and electrical grids (which they were first invented for, as mentioned above)
- Approximate NP-complete graph optimization (e.g., travel salesman problem)
- Image analysis

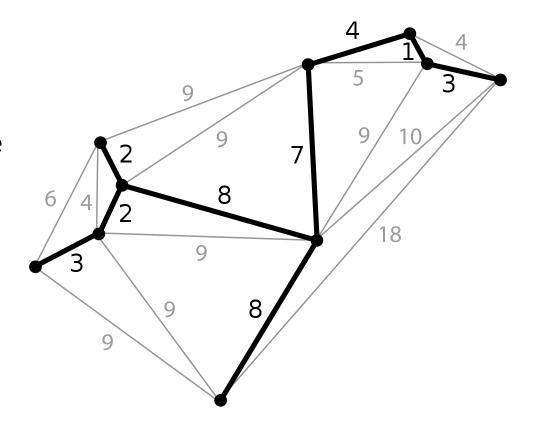


Image from Wikipedia



Minimum Spanning Tree Problem

- Two efficient greedy algorithms presented here: Prim's and Kruskal's.
- Each selects edges in order of increasing weight but avoids creating a cycle.

Prim's

- Prim's algorithm maintains a tree at each stage that grows to span
- Prim's implementation very similar to Dijkstra, runs in time $O((m+n)\log n)$

Kruskal's

- Kruskal maintains a forest whose trees are combined into one spanning tree.
- Kruskal can be implemented to run in time $O(m \log n)$.



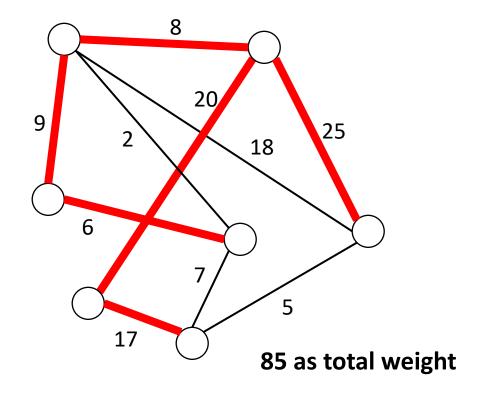
Finding Spanning Trees

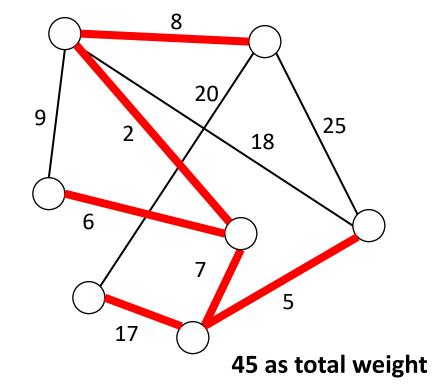
- Finding a spanning tree is easy.
- Steo1: Run DFS or BFS from any of the vertices.
- Step2: If G is connected, there will be only one search tree and this will be a spanning tree.



Spanning Trees in Weighted Graphs (Contd.)

The total weight of a spanning tree is the sum of the weights of its edges.

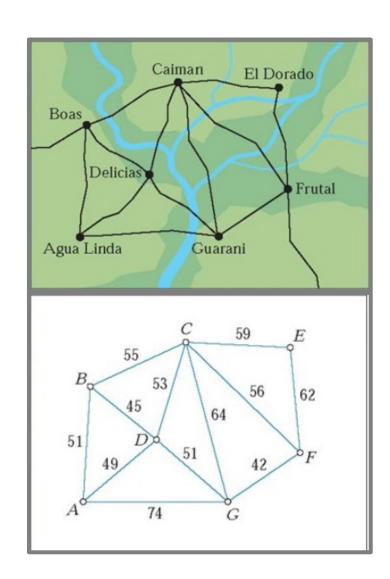






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Prim's Algorithm

- Prim's algorithm is a modified version of Dijkstra's.
- We simply pick one of the vertices in *G* as the source.
- We need to keep track of the edges used to reach a vertex.

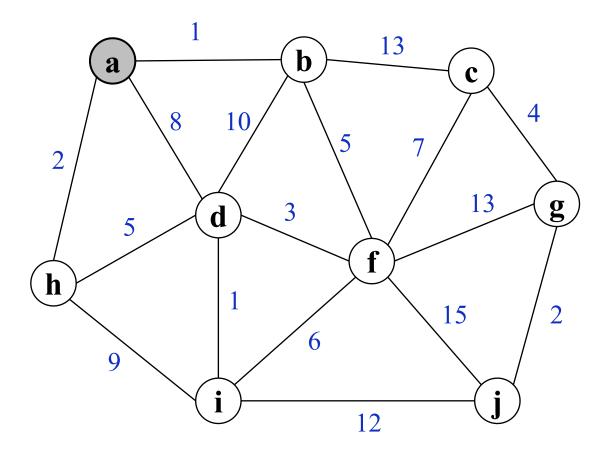
Algorithm 1 Prim's algorithm.



```
1: function PRIM(weighted graph(G, c); vertex s \in V(G))
2:
           priority queue Q
            array colour[0..n-1], pred[0..n-1]
3:
4:
           for u \in V(G) do
5:
                 colour[u] \leftarrow WHITE; pred[u] \leftarrow null
            colour[s] \leftarrow \mathsf{GREY}
6:
7:
            Q.insert(s, 0)
8:
            while not Q. isEmpty() do
9:
                 u \leftarrow Q.peek()
10:
                 for each x adjacent to u do
11:
                      t \leftarrow c(u, x)
12:
                      if colour[x] = WHITE then
                           colour[x] \leftarrow GREY; pred[x] \leftarrow u
13:
14:
                           Q.insert(x,t)
                      else if colour[x] = GREY and Q.getKey(x) > t then
15:
16:
                           Q.decreaseKey(x,t); pred[x] \leftarrow u
17:
                 Q.delete()
18:
                 colour[u] \leftarrow BLACK
19:
            return pred
```



Priority Queue Q: (a, 0)



Pred:
a: null
b: null
c: null
d: null
f: null

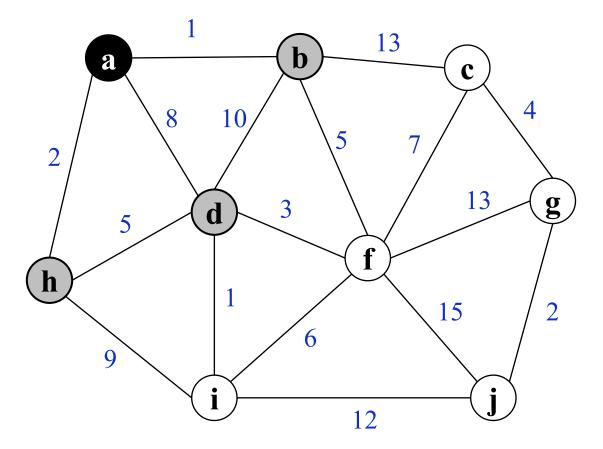


Priority Queue Q:

(b, 1)

(d, 8)

(h, 2)



Pred:

a: null

b: a

c: null

d: a

f: null

g: null

h: a

i: null

j: null



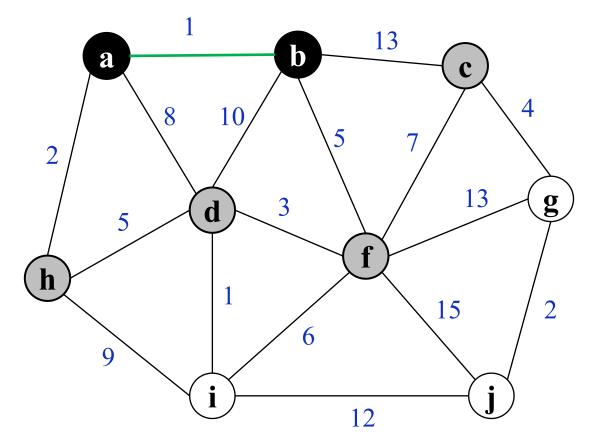
Priority Queue Q:

(d, 8)

(h, 2)

(c, 13)

(f, 5)



Pred:

a: null

b: a

c: b

d: a

f: b

g: null

h: a

i: null

j: null



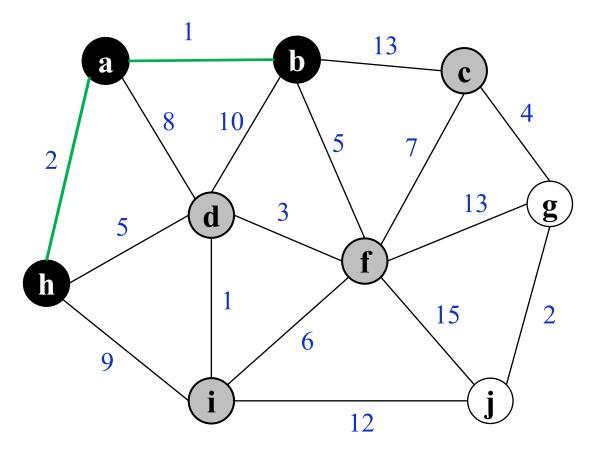
Priority Queue Q:

(d, 5)

(c, 13)

(f, 5)

(i, 9)



Pred:

a: null

b: a

c: b

d: h

f: b

g: null

h: a

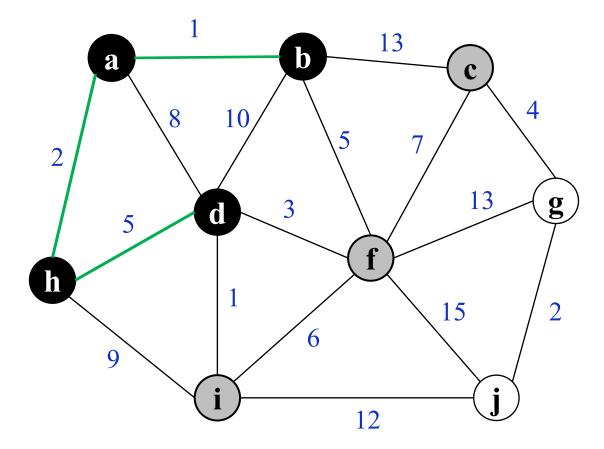
i: h

j: null



Priority Queue Q: (c, 13) (f, 3)

(i, 1)



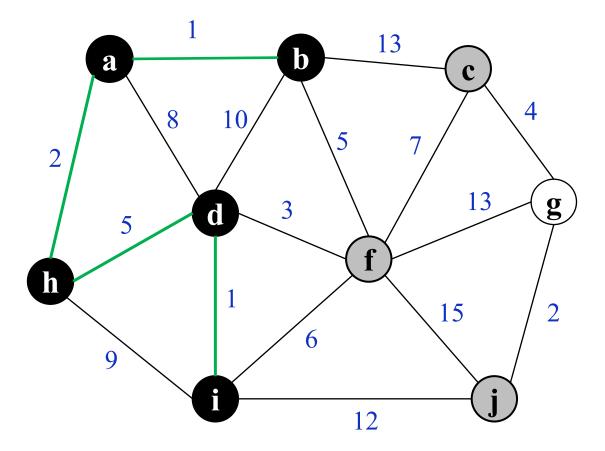
Pred:
a: null
b: a
c: b
d: h
f: d
g: null
h: a
i: d
j: null



Priority Queue Q: (c, 13)

(f, 3)

(j, 12)



Pred:

a: null

b: a

c: b

d: h

f: d

g: null

h: a

i: d

j: i

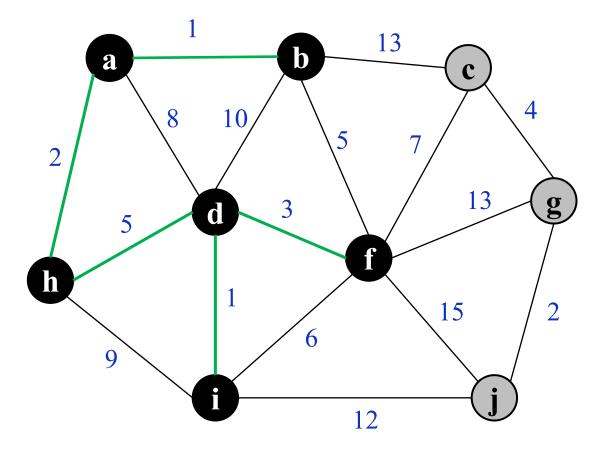


Priority Queue Q:

(c, 7)

(j, 12)

(g, 13)



Pred:

a: null

b: a

c: f

d: h

f: d

g: f

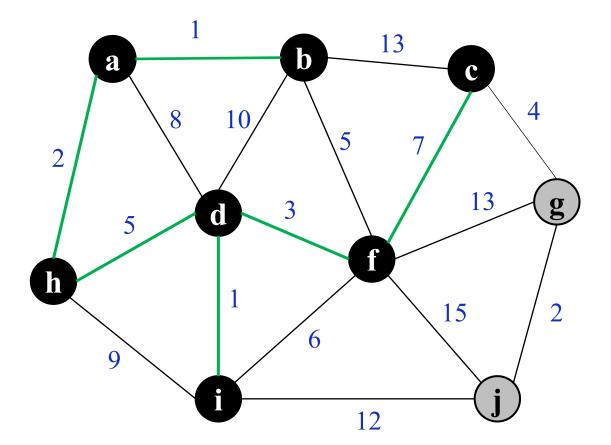
h: a

i: d

j: i



Priority Queue Q: (j, 12) (g, 4)

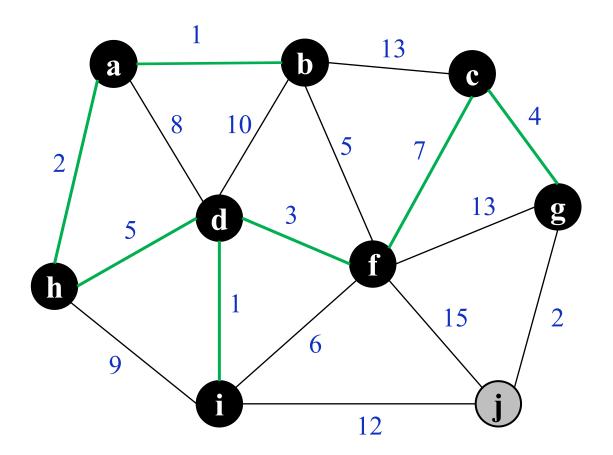


Pred:
a: null
b: a
c: f
d: h
f: d
g: c
h: a
i: d

j: i



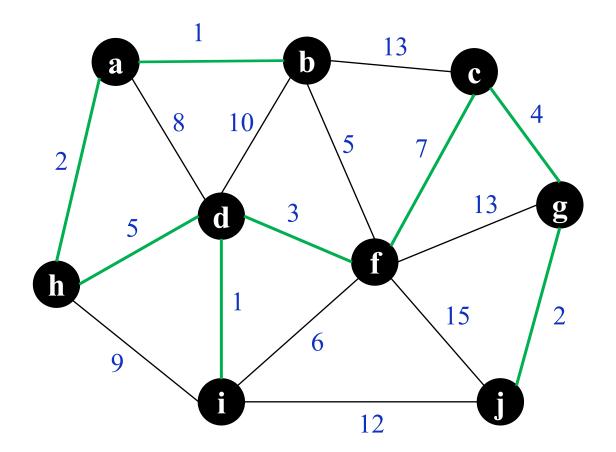
Priority Queue Q: (j, 2)



Pred:
a: null
b: a
c: f
d: h
f: d
g: c
h: a
i: d
j: g



Priority Queue Q:



Pred:

a: null

b: a

c: f

d: h

f: d

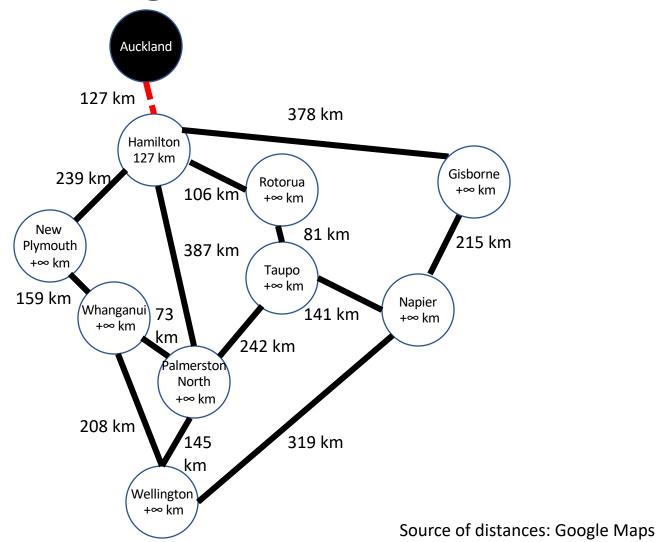
g: c

h: a

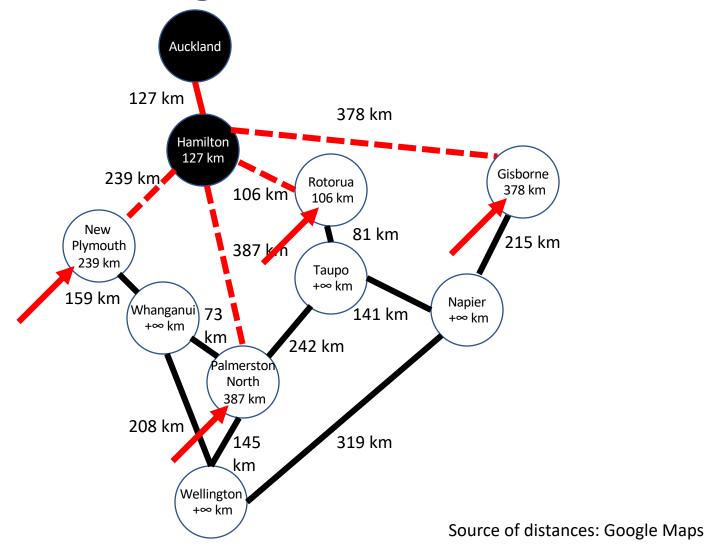
i: d

j: g

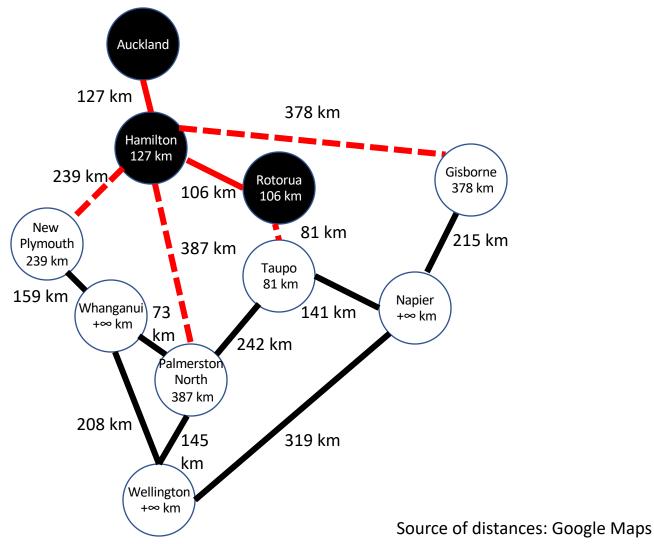




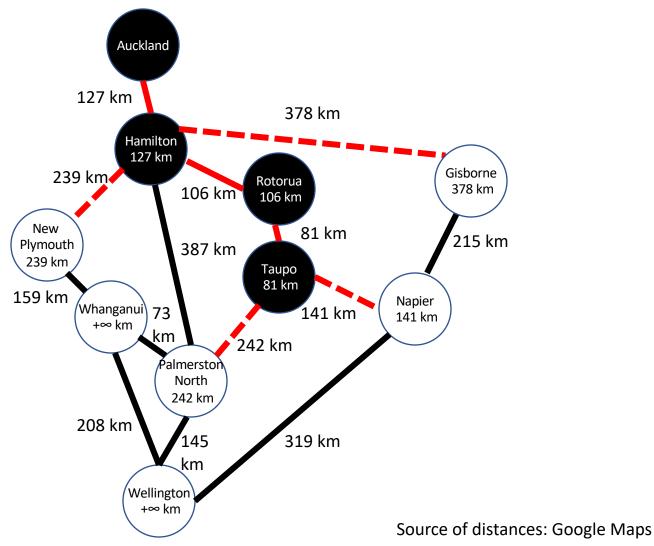




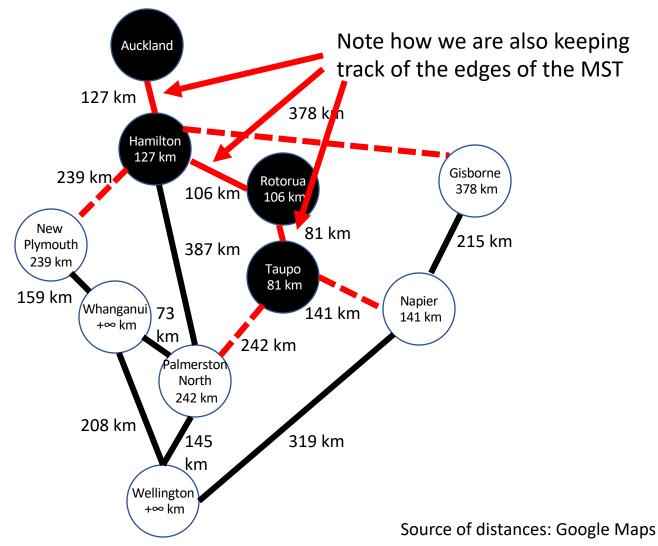




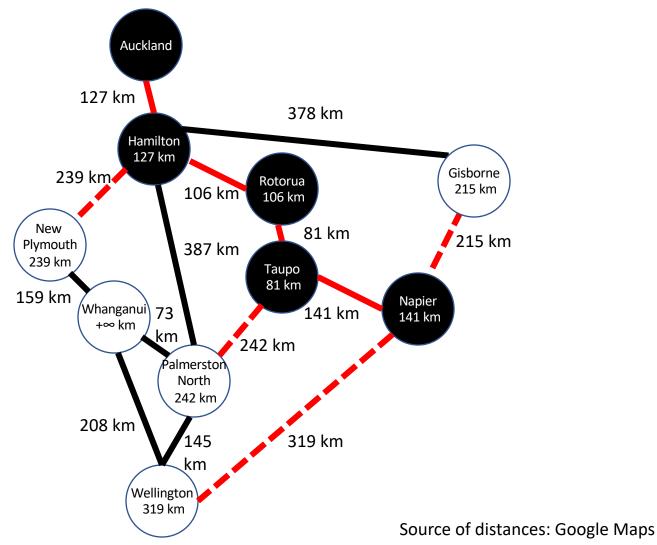




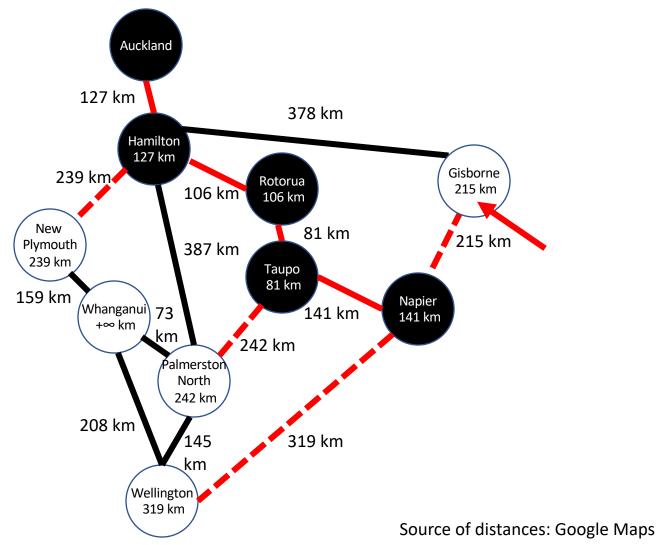




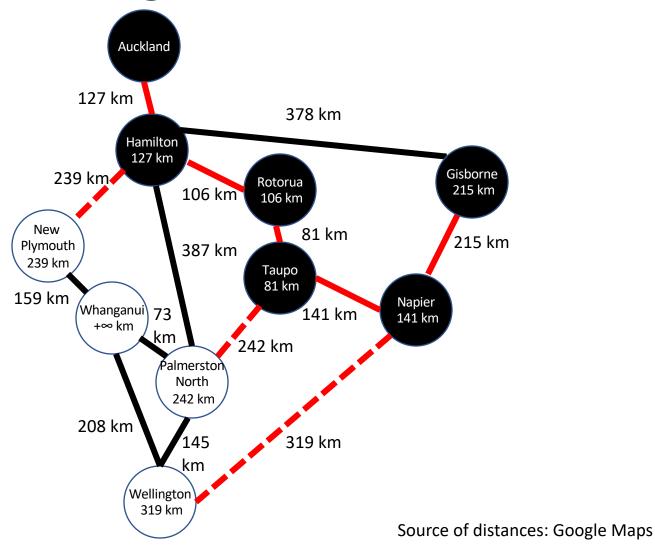




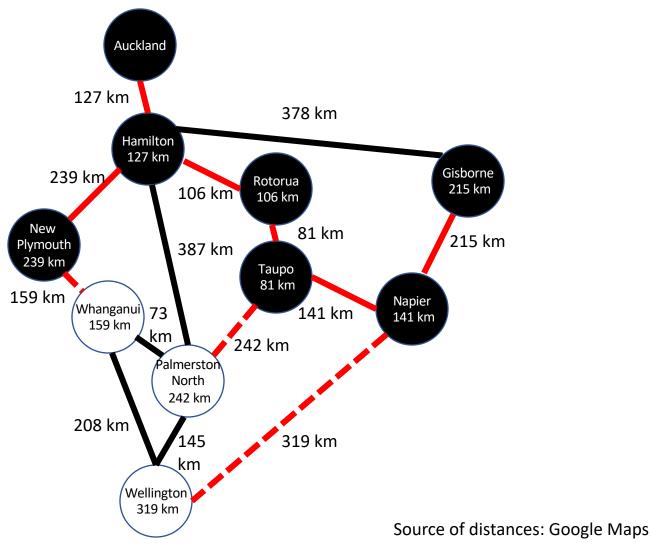




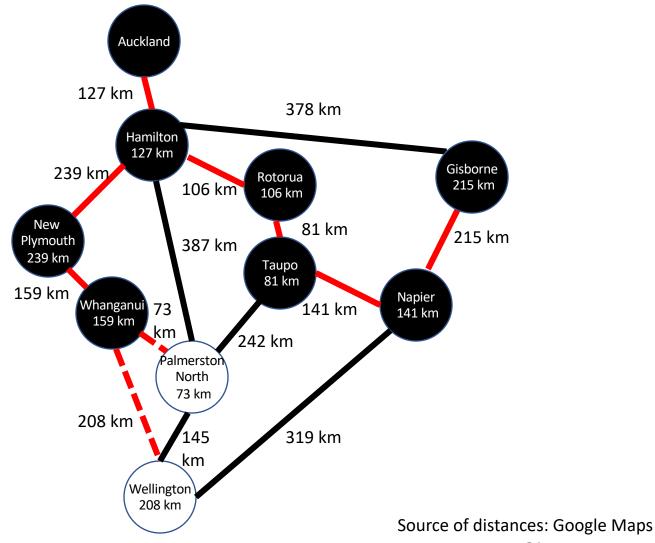




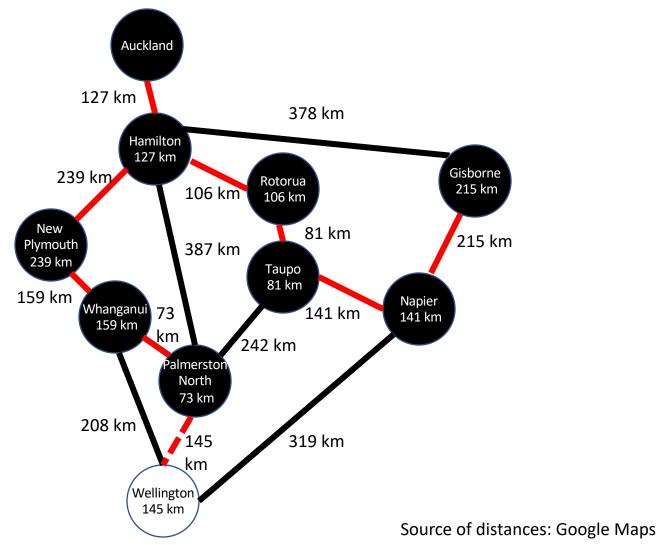




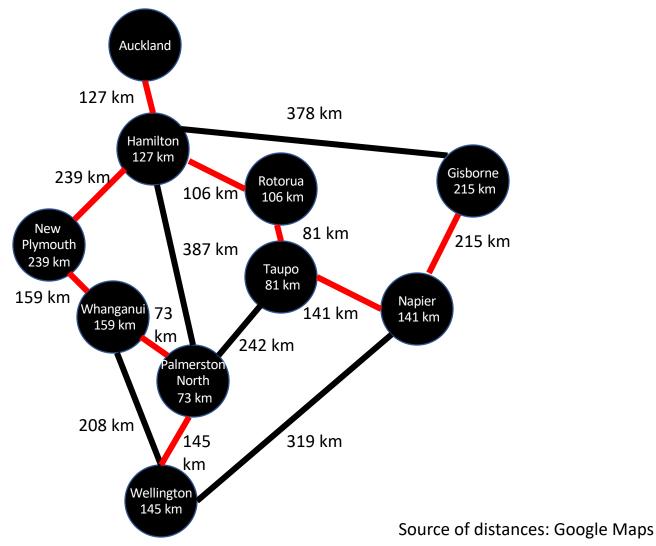














Time Complexity Analysis

- Prim's algorithm is essentially the same as Dijkstra's
- The time complexity therefore is the same as Dijkstra's.



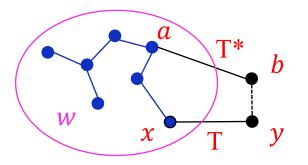
Correctness of Prim's Algorithm

- Proof by contradiction.
- Let T be a spanning tree that is obtained by Prim's algorithm and assume that T is not of minimum weight. Let $e_1, e_2, \ldots, e_{n-1}$ be the edges of T chosen in order. Let T^* be an MST with edges $f_1, f_2, \ldots, f_{n-1}$.
- We may assume that T^* is chosen such that $i \in \{1,2,...,n-1\}$ with $e_j = f_j$ for all j < i is maximum over all choices for T^* . Let w be the set of vertices selected in T before e_i is selected, and (x, y) be the first edge in T but not in T^* . Let $E = E(T^*) \cup \{(x, y) (a, b)\}$ be the edge set of a spanning tree.



Correctness of Prim's Algorithm (Contd.)

- Case 1: c(a,b)>c(x,y). Then E has a smaller weight than T*.
- Case 2: c(a,b) < c(x,y). Then the algorithm chooses (a,b) and not (x,y).
- Case 3: c(a,b)=c(x,y). Then there exists an MST whose first i edges are identical to T.

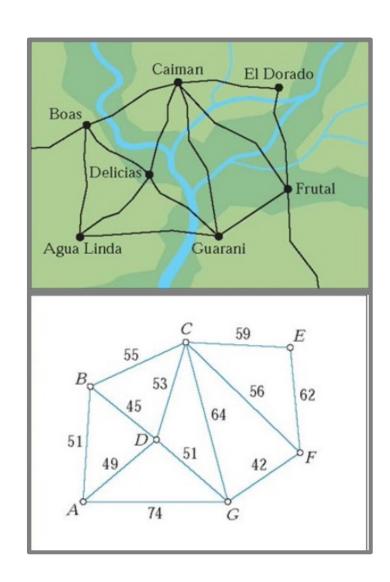




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Kruskal's Algorithm

- Prim's builds an MST using black vertices
- Kruskal's builds a forest that then converges to a MST
- Both algorithms build the tree avoiding to add cycles



Kruskal's Algorithm (Contd.)

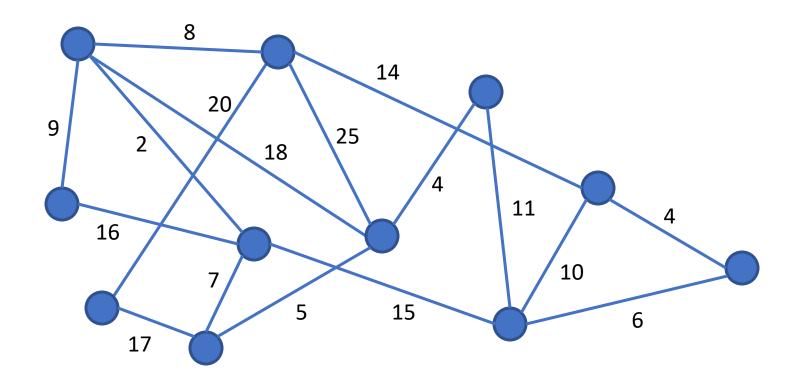
- Step1: First sort the edges by their weights
- Step2: Add an edge to the forest if the edge does not connect two vertices already in the same tree



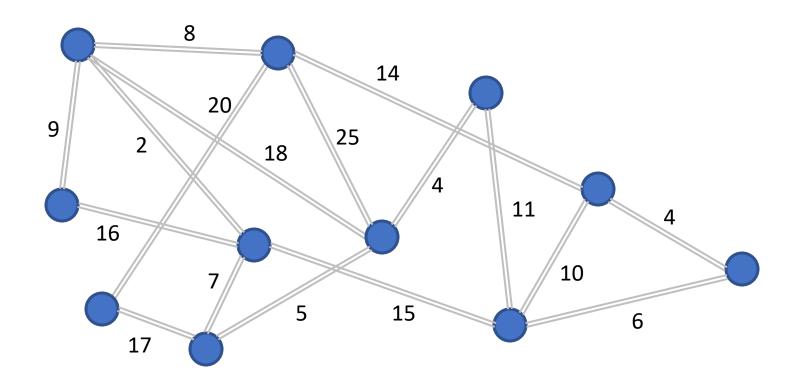
Kruskal's Algorithm

Algorithm 2 Kruskal's algorithm. 1: **function** KRUSKAL(weighted digraph(G, c)) 2: disjoint sets ADT A initialize A with each vertex in its own set 3: sort the edges in increasing order of cost 4: 5: for each edge $\{u, v\}$ in increasing cost order do if not A. set(u) = A. set(v) then 6: add this edge 7: A.union(A.set(u), A.set(v))8: 9: return A

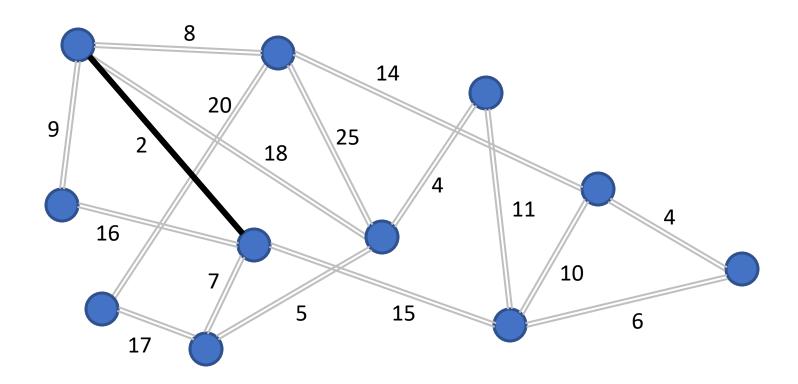






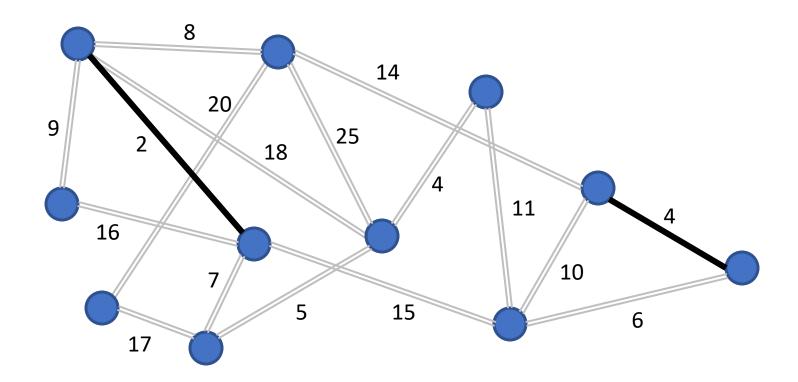






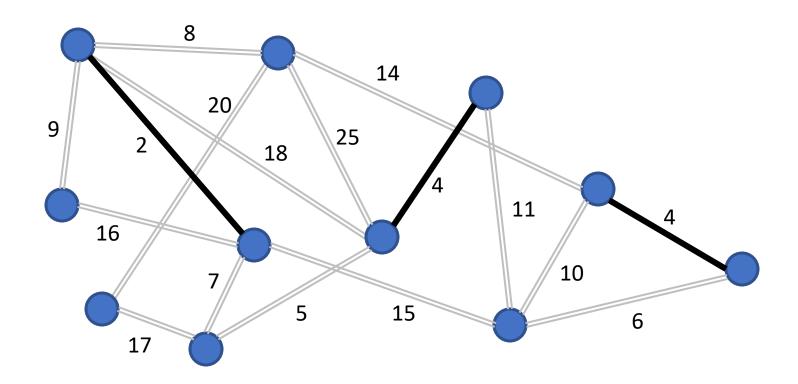






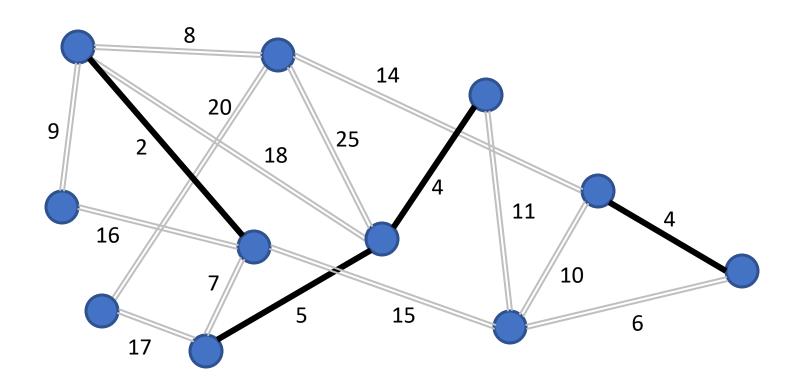
$$i=1$$





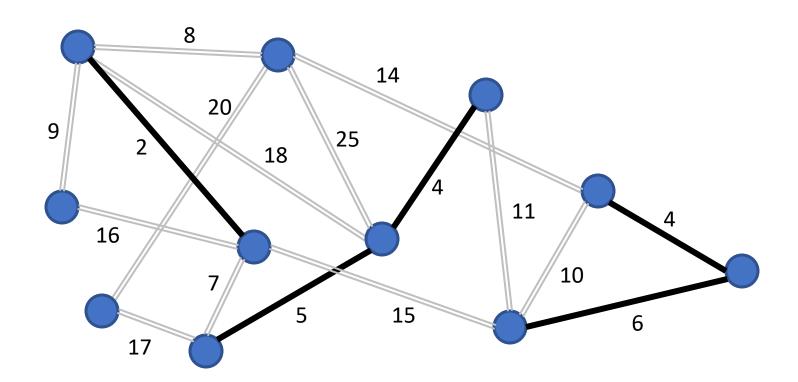






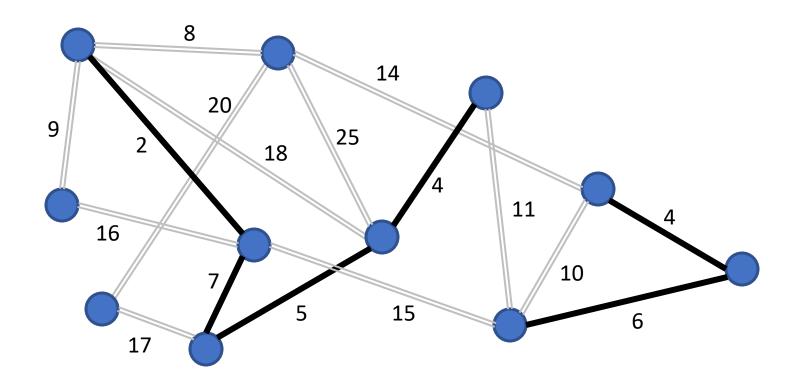






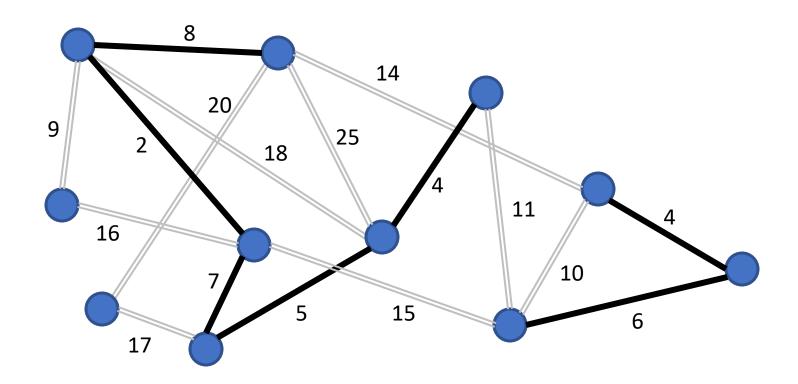






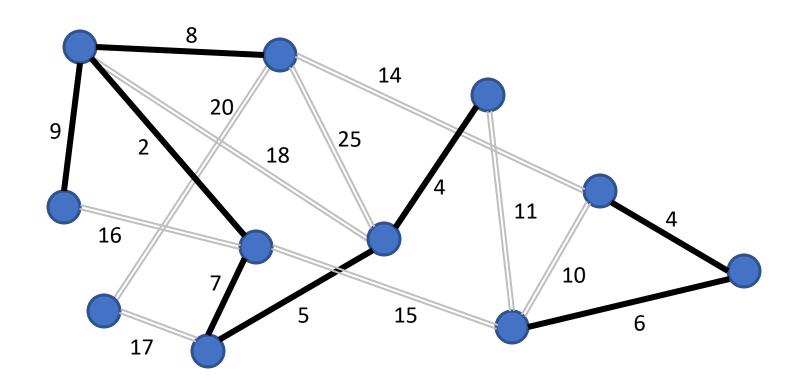






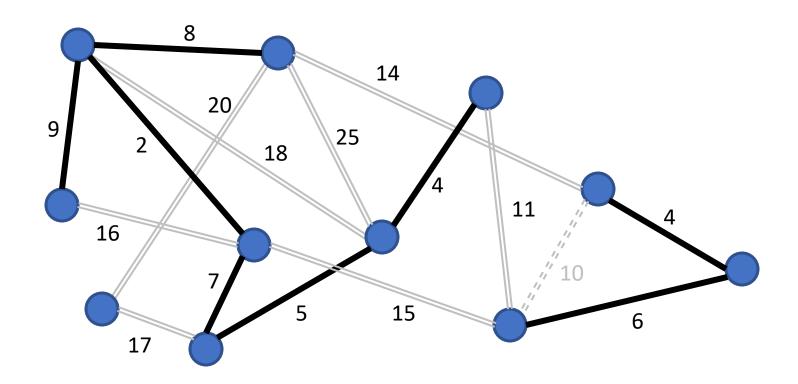
$$i=6$$





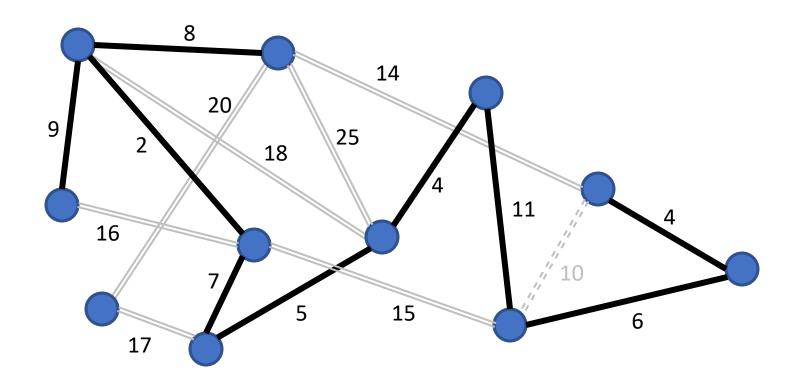






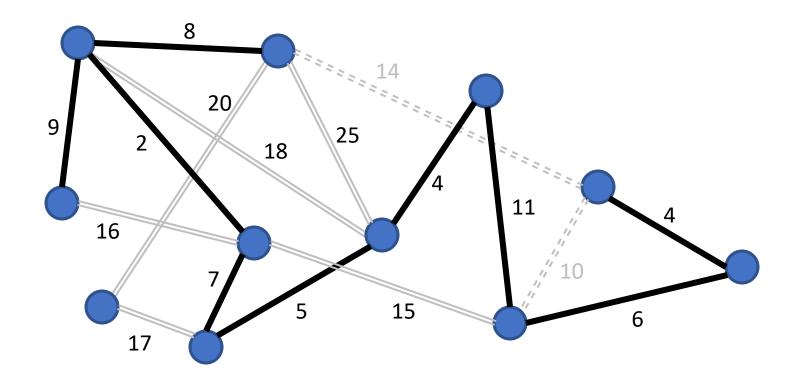






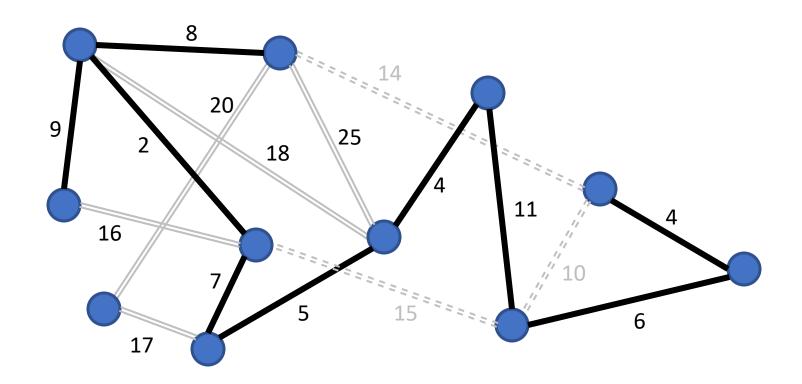






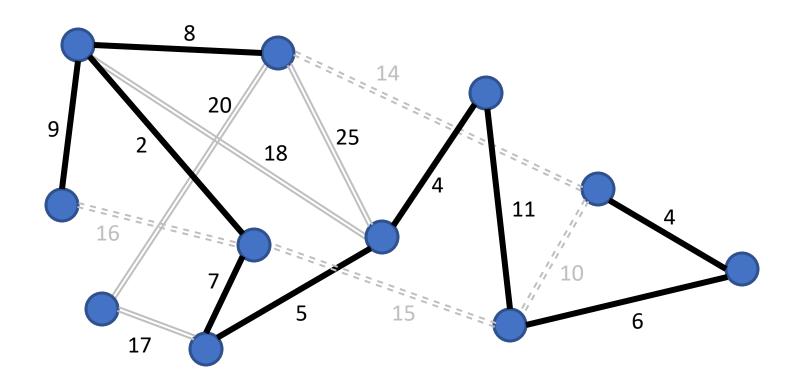
$$i = 10$$





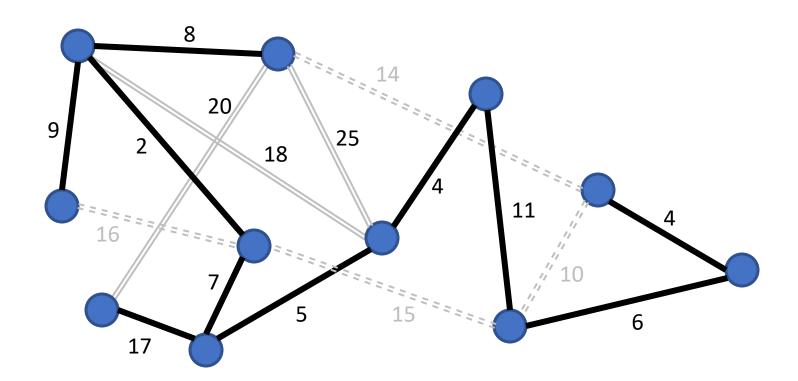
i = 11





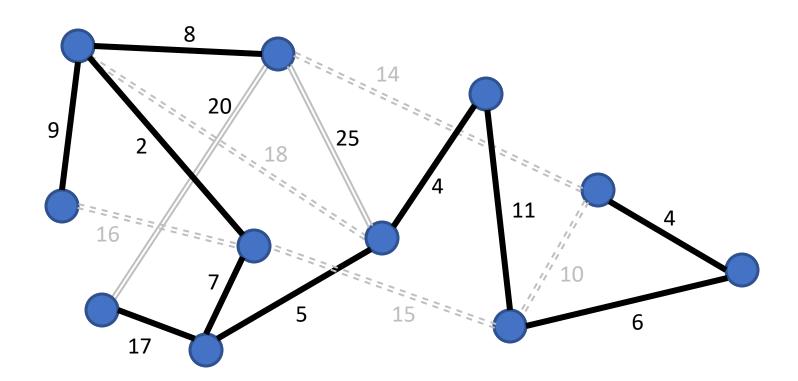
$$i=12$$





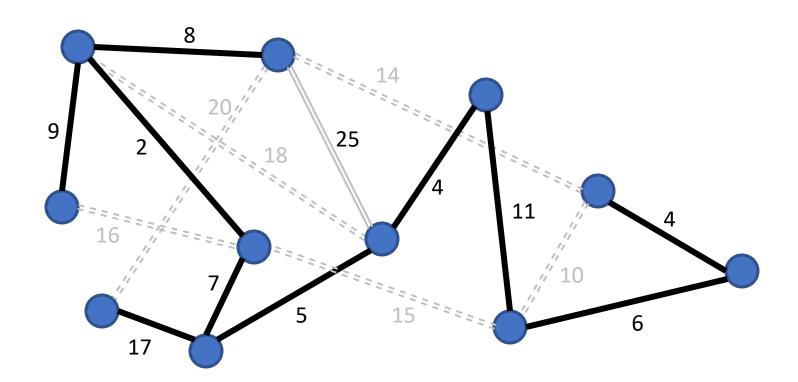
$$i = 13$$





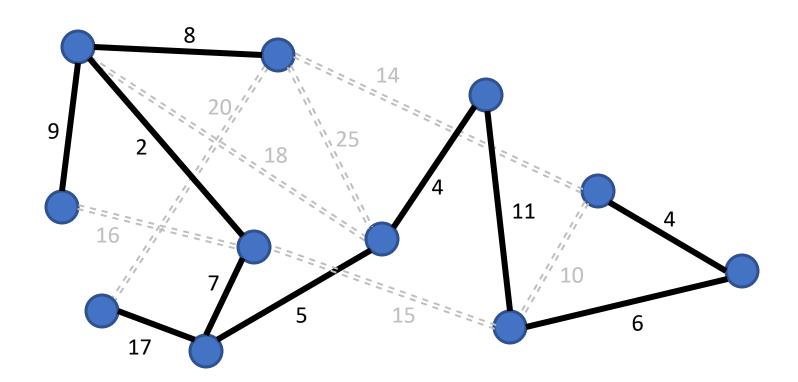
$$i = 14$$





$$i = 15$$





$$i = 16$$



Disjoint Set ADT

- Disjoint Sets ADT: Used to represent a collection of sets containing objects that are related to each other. Two standard operations:
 - Union operation merges two sets by names (denoted by one of the objects in the set).
 - Find operation given an object, returns the set (with its name) the object belongs to

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Example:
```

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Initial Sets = \{1,4,8\}, \{2,3\}, \{6\}, \{7\}, \{5,9,10\}
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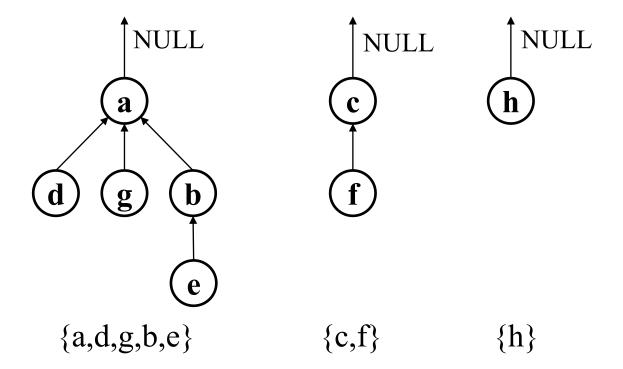
Find(4) returns a set named 1

Union(2, 6) results in a new set $\{2,3,6\}$ with name 2.



Up-Tree Data Structure

- Each disjoint set is an up-tree with its root as its representative member
- All members of a given set are nodes in that set's up-tree





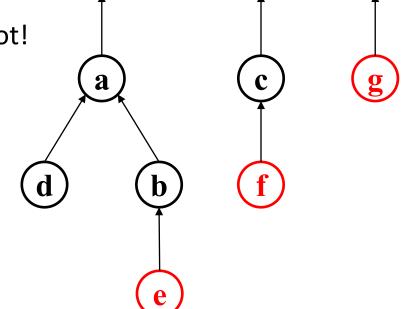
Example: Find

• Find: Just follow parent pointers to the root!

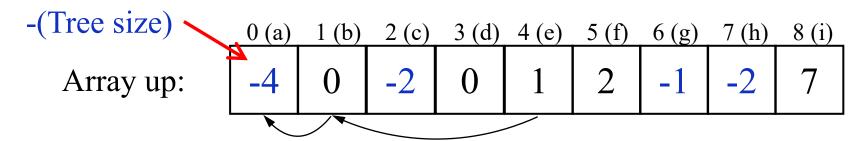
$$find(f) = c$$

$$find(e) = a$$

$$find(g) = g$$



Runtime depends on tree depth





Example: Union

• Union: put root under the root of the other tree

union(c,a)

Runtime = O(1)

e
3 (d) 4 (e) 5 (f) 6 (g) 7 (h) 8 (i)
0 1 2 -1 -2 7

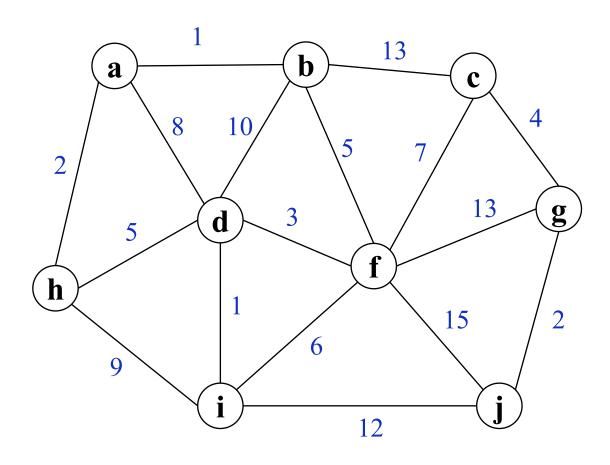
Array up:

 0 (a)
 1 (b)
 2 (c)
 3 (d)
 4 (e)
 5 (f)
 6 (g)
 7 (h)
 8 (i)

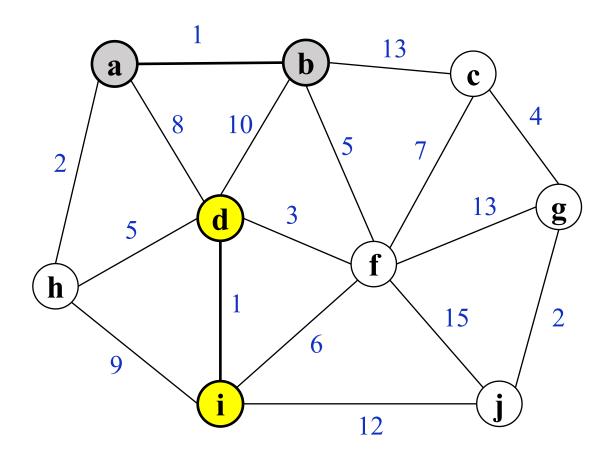
 2
 0
 -6
 0
 1
 2
 -1
 -2
 7

Change a (from -4) to c(=2), update size

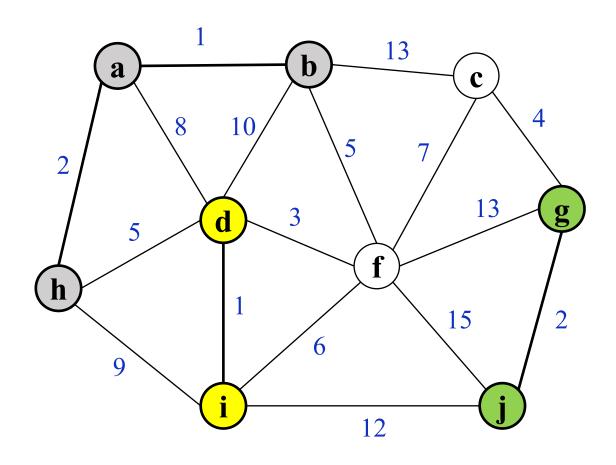




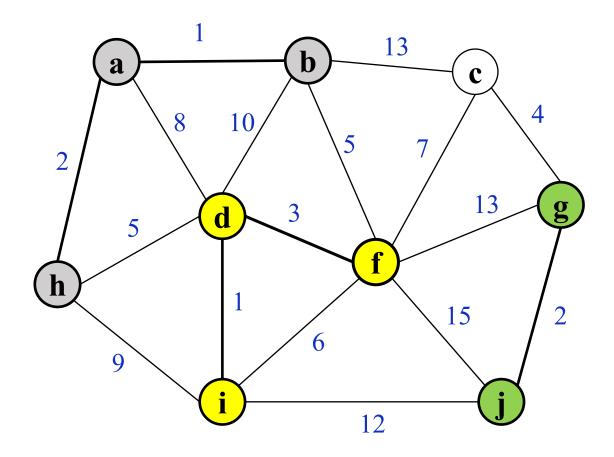




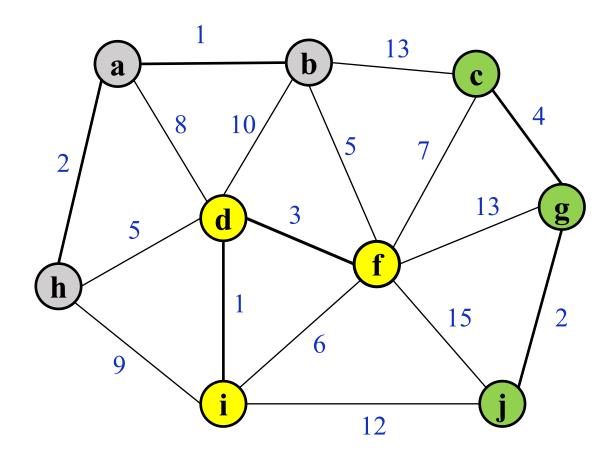




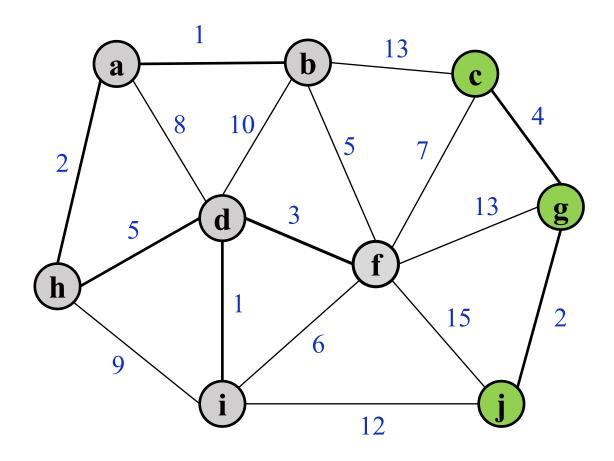




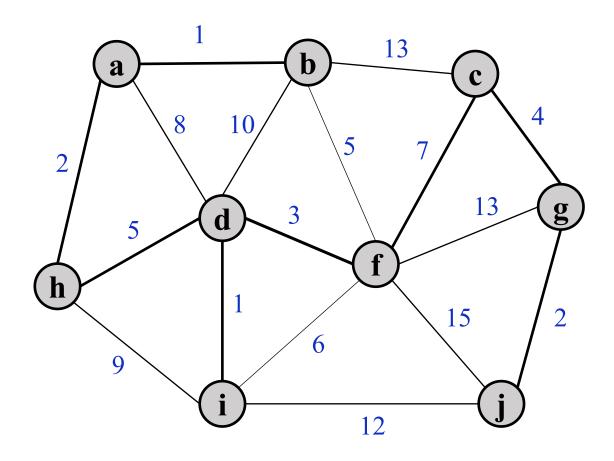














Time Complexity Analysis

- Time complexity is dominated by having to sort the edges by weight: $O(e \log e)$, the time to sort.
 - Note e is at most n^2 and $\log e = \log n^2 = 2 \log n$.
 - Therefore $O(e \log n)$
- Kruskal's algorithm is a little harder to program but easier to do by hand



Minimum Spanning Tree (MST) Summary

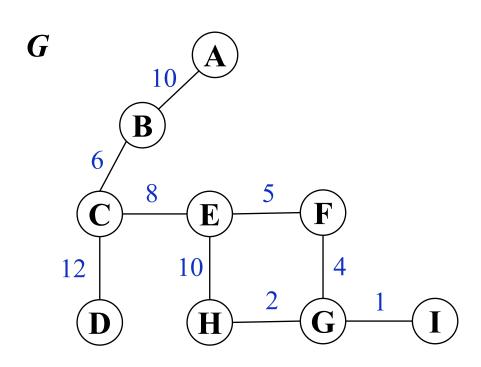
• Fact.

- 1. The most expensive edge, if unique, of a cycle in an edge-weighted graph G is not in any MST. (Otherwise, at least one of those equally expensive edges of the cycle must not be in each MST.)
- 2. The minimum cost edge, if unique, between any non-empty strict subset S of V(G) and the $V(G)\$ S is in the MST. (Otherwise, at least one of these minimum cost edges is in each MST.)

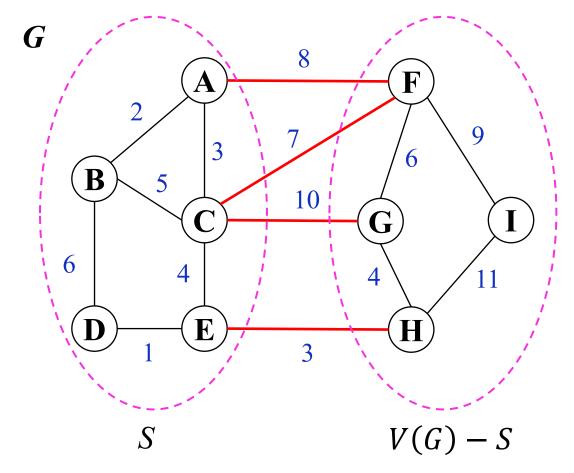


Max/Min Cost Edges in MST

 $S = \{A, B, C, D, E\}$ $V(G) - S = \{F, G, H, I\}$ $MST \text{ of } G \text{ contains } \{E, H\}$



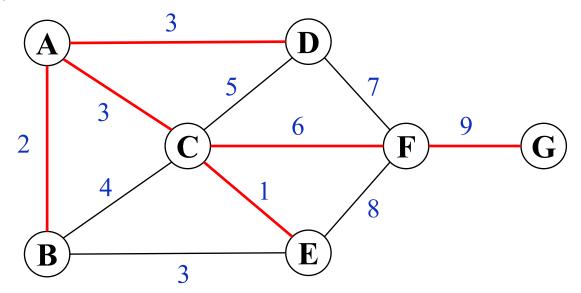
MST does not contain $\{E, H\}$





Prim's Algorithm



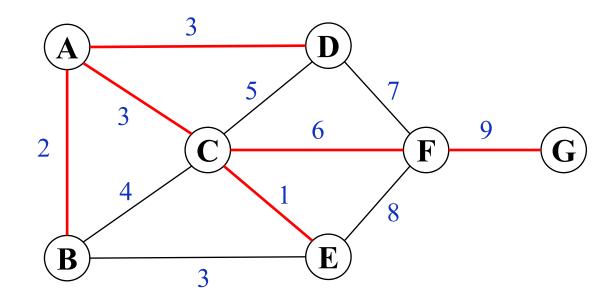


 ${A,B},{A,C},{C,E},{A,D},{C,F},{F,G}$

MST of weight: 24



Kruskal's Algorithm



 $\{C, E\}, \{A, B\}, \{A, C\}, \{A, D\}, \{C, F\}, \{F, G\}$ 1 2 3 3 6 9 MST of weight: 24



Comparing the Prim's and Kruskal's Algorithms

Both algorithms choose and add at each step a min-weight edge from the remaining edges, subject to constraints

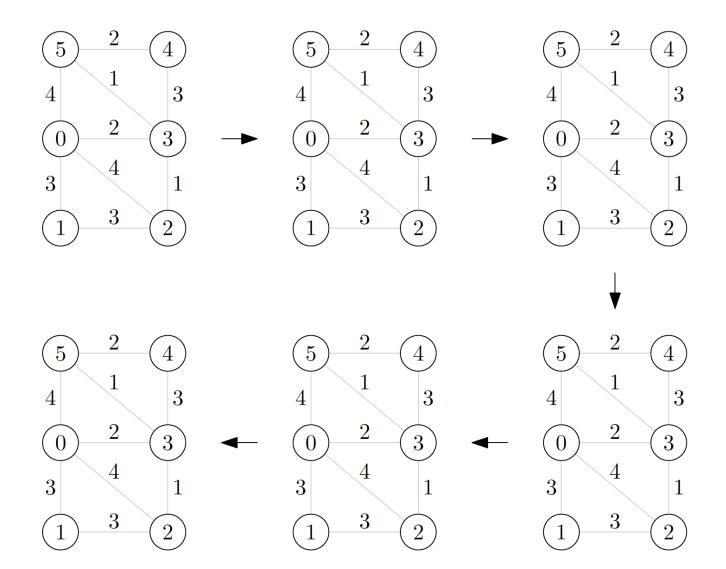
Prim's MST algorithm:

- Start at a root vertex.
- Two rules for a new edge:
 - 1. No cycle in the subgraph built so far.
 - 2. Connect the subgraph built so far.
- Terminate if no more edges to add can be found.
- At each step: an acyclic connected subgraph being a tree.

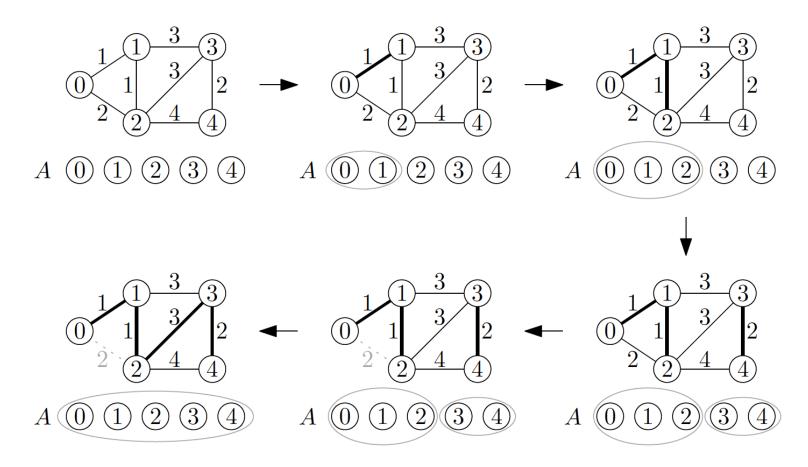
Kruskal's MST algorithm:

- Start at a min-weight edge.
- One rule for a new edge:
 - No cycle in a forest of trees built so far.
- Terminate if no more edges to add can be found.
- At each step: a forest of trees merging as the algorithm progresses (can find a spanning forest for a disconnected graph).

Example 32.4. Execute Prim's algorithm



Example 32.6. Application of Kruskal's algorithm on a graph shown until an MST is found. Note that the edge {0, 2} with weight 2 is not added, because 0 and 2 are already in the same set in A.





SUMMARY

- Terminology
 - Spanning Trees
 - Minimum Spanning Trees (MST)
- Finding MST es Algorithms
 - Prim's Algorithm
 - Kruskal's Algorithm
 - Time Complexity Analysis
 - Comparison

