### CSE 373: Minimum Spanning Trees: Prim and Kruskal

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### Minimum spanning trees

Punchline: a MST of a graph connects all the vertices together while minimizing the number of edges used (and their weights). Minimum spanning trees

Given a connected, undirected graph G = (V, E), a minimum spanning tree is a subgraph G' = (V', E') such that...

V = V' (G' is spanning)

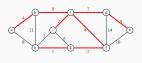
- There exists a path from any vertex to any other one
  - ➤ The sum of the edge weights in E' is minimized.

### In order for a graph to have a MST, the graph must...

- ▶ ...be connected there is a path from a vertex to any other
  - vertex. (Note: this means  $|V| \le |E|$ ).
- be undirected

## Minimum spanning trees: example

An example of an minimum spanning tree (MST):



### Minimum spanning trees: Applications

### Example questions:

- ➤ We want to connect phone lines to houses, but laving down cable is expensive. How can we minimize the amount of wire we must install?
- ▶ We have items on a circuit we want to be "electrically equivalent". How can we connect them together using a minimum amount of wire?

### Other applications:

- ► Implement efficient multiple constant multiplication
- Minimizing number of packets transmitted across a network
- ► Machine learning (e.g. real-time face verification)
- ▶ Graphics (e.g. image segmentation)

### Minimum spanning trees: properties

### Important properties:

- ► A valid MST cannot contain a cycle vertices are no longer connected.
- ▶ If we add or remove an edge from an MST, it's no longer a valid MST for that graph. Adding an edge introduces a cycle; removing an edge means
- ▶ If there are |V| vertices, the MST contains exactly |V| 1 edges
- ▶ An MST is always a tree.
- ▶ If every edge has a unique weight, there exists a unique MST.

## Minimum spanning trees: algorithm

Design question: how would you implement an algorithm to find the MST of some graph, assuming the edges all have the same weight?

### Hinte-

- ► Try modifying DFS or BFS.
- Try using an incremental approach: start with an empty graph, and steadily add nodes and edges.

### Minimum spanning trees: approach 1, adding nodes

Intuition: We start with an "empty" MST, and steadily grow it.

Core algorithm:

- 1. Start with an arbitrary node.
- 2. Run either DFS or BFS, storing edges in our stack or queue.
- 3. As we visit nodes, add each edge we remove to our MST.

### Minimum spanning trees: approach 1, adding nodes

An example using a modified version of DFS:



Stack:

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# Minimum spanning trees: approach 1, adding nodes

An example using a modified version of DFS:



Stack: (a, b), (a, d),

Minimum spanning trees: approach 1, adding nodes

An example using a modified version of DFS:



Stack: (a, b), (d, e), (d, f), (d, g),

# Minimum spanning trees: approach 1, adding nodes

An example using a modified version of DFS:



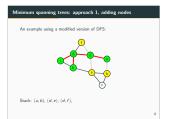
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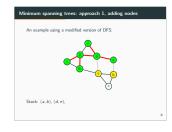
Minimum spanning trees: approach 1, adding nodes

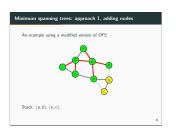
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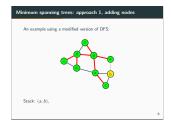


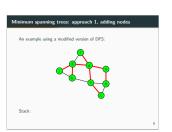
Stack: (a, b), (d, e), (d, f), (g, h),













### Interlude: finding the shortest path

Review: How do we find the shortest path between two vertices?

- ► If the graph is unweighted: run BFS
- ► If the graph is weighted: run Diikstra's

How does Diikstra's algorithm work?

- Give each vertex v a "cost": the cost of the shortest-known path so far between v and the start.
   (The cost of a path is the sum of the edge weights in that path)
- Pick the node with the smallest cost, update adjacent node costs, repeat

### Minimum spanning trees: approach 1, adding nodes

Intuition: We can use the same idea to find a MST!

Core idea: Use the exact same algorithm as Dijkstra's algorithm, but redefine the cost:

### ► Previously, for Diikstra's:

The cost of vertex v is the cost of the shortest-known path so far between v and the start

### ► Now:

The cost of vertex v is the cost of the shortest-known path so far between v and any node we've visited so far

This algorithm is known as Prim's algorithm.

### Compare and contrast: Dijkstra vs Prim

### Pseudocode for Dijkstra's algorithm:

def dijkstra(start):

backpointers = new SomeDictionary(Vertex, Vertex>()

for (v : vertices):

set cost(v) to infinity set cost(start) to 0

while (we still have unvisited nodes) current - get next smallest node

for (edge : current.getOutEdges()):
 newCost = min(cost(current) + edge.cost, cost(edge.dst))
 update cost(edge.dst) to newCost
 backcostners.cut(edge.dst.) edge.arc)

return backpointer

..

### Compare and contrast: Dijkstra vs Prim

### Pseudocode for Prim's algorithm:

def prim(start):
 backpointers = new SomeDictionaryVertex>()

for (v : vertices):

set cost(v) to infinity

while (we still have unvisited nodes): current = get next smallest node

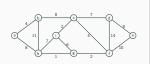
for (edge : current.getOutEdges()):

newCost = min(edge.cost, cost(edge.dst))
update cost(edge.dst) to newCost
backpointers.put(edge.dst, edge.src)

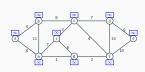
return backpointers

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### Prim's algorithm: an example

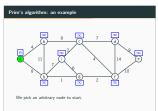


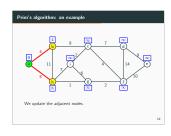
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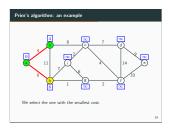


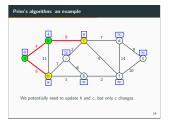
We initially set all costs to  $\infty$ , just like with Dijkstra.

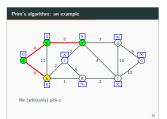
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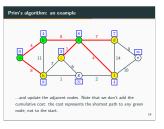


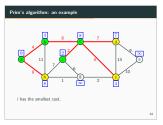


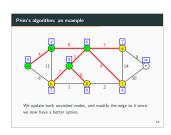


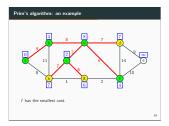


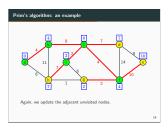


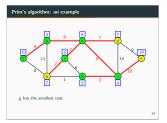


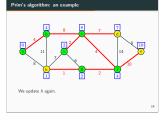


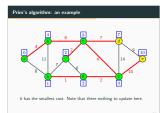


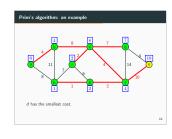


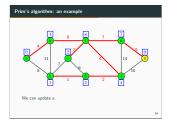


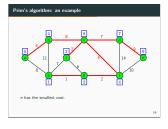


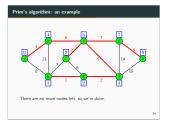


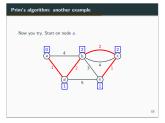












### Analyzing Prim's algorithm

Question: What is the worst-case asymptotic runtime of Prim's algorithm?

Answer: The same as Dijkstra's:  $O(|V|t_x + |E|t_u)$  where...

- $\blacktriangleright t_{\rm s} = {\rm time}$  needed to get next smallest node
- ▶ t<sub>u</sub> = time needed to update vertex costs

So,  $\mathcal{O}\left(|V|\log(|V|) + |E|\log(|V|)\right)$  if we stick to data structures we know how to implement;  $\mathcal{O}\left(|V|\log(|V|) + |E|\right)$  if we use Fibonacci heaps.

### Minimum spanning trees, approach 2

Recap: Prim's algorithm works similarly to Dijkstra's – we start with a single node, and "grow" our MST.

A second approach: instead of "growing" our MST, we...

- Initially place each node into its own MST of size 1 so, we start with |V| MSTs in total.
- Steadily combine together different MSTs until we have just one left
- How? Loop through every single edge, see if we can use it to join two different MSTs together.

This algorithm is called Kruskal's algorithm

### Kruskal's algorithm

An example, for unweighted graphs. Note: each MST has a different color.



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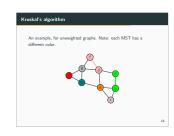
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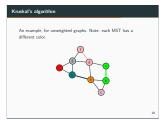
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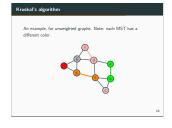


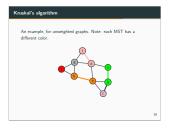
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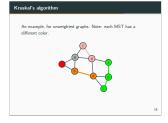
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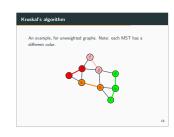


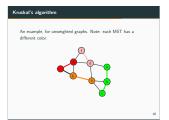


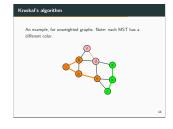


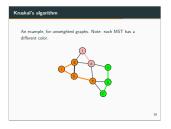


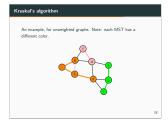
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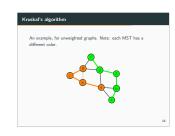




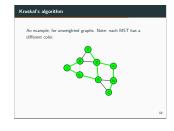


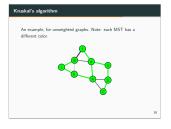


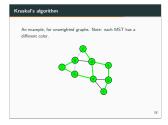
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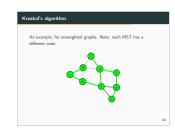
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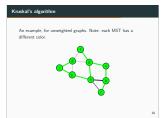


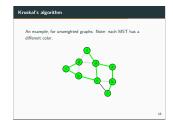


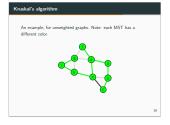


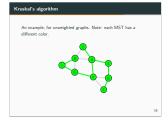
# Kruskal's algorithm An example, for unweighted graphs. Note: each MST has a different color.











### Kruskal's algorithm: weighted graphs

Question: How do we handle edge weights?

Answer: Consider edges sorted in ascending order by weight.

So, we look at the edge with the smallest weight first, the edge with the second smallest weight next, etc.

### Kruskal's algorithm: pseudocode

### Pseudocode for Kruskal's algorithm:

def kruskal():

mst - new SomeSet<Edge>()

for (v : vertices):
 makeMST(v)

sort edges in ascending order by their weight

for (edge : edges):
 if findMST(edge.src) != findMST(edge.dst):
 union(edge.src, edge.dst)
 mst.add(edge)

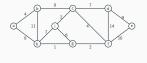
► makeMST(v): stores v as a MST containing just one node

- ► findMST(v): finds the MST that vertex is a part of
- vertices, using the edge (u,v)

▶ union(u, v): combines the two MSTs of the two given

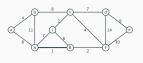
## Kruskal's algorithm: example with a weighted graph

Now you try:



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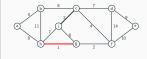
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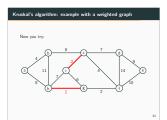
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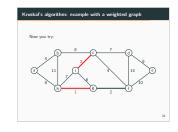


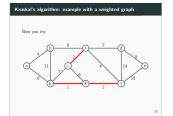
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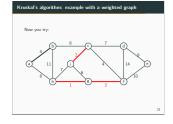
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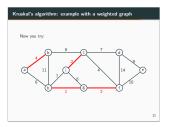


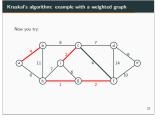


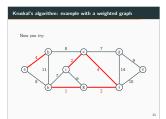


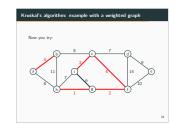


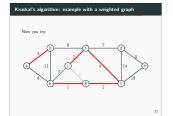


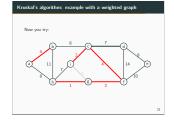


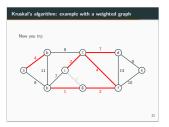


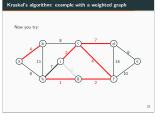


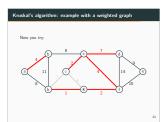


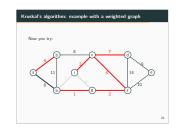


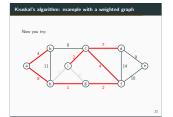


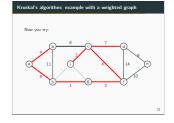


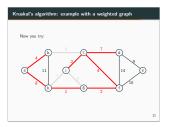


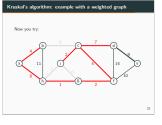


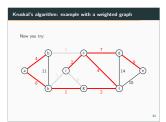


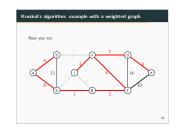


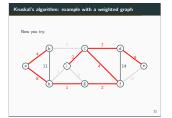


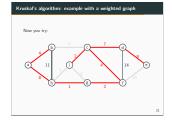


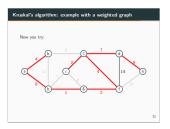


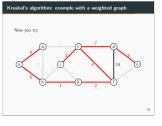












### Kruskal's algorithm: example with a weighted graph





### Kruskal's algorithm: analysis

### What is the worst-case runtime?

def kruskal(): mst - new SomeSet<Edge>()

for (v : vertices):
 makeMST(v)

sort edges in ascending order by their weight

for (edge : edges):
 if findMST(edge.src) != findMST(edge.dst):
 union(edge.src, edge.dst)
 mst.add(edge)

# Note: assume that...

- ▶ makeMST(v) takes O(t<sub>m</sub>) time
- ▶ findMST(v): takes O(t<sub>ℓ</sub>) time
- ▶ union(u, v): takes O(t<sub>u</sub>) time

### Kruskal's algorithm: analysis

- ► Making the |V| MSTs takes O(|V|-t<sub>m</sub>) time
- ▶ Sorting the edges takes  $O(|E| \cdot \log(|E|))$  time, assuming we use a general-purpose comparison sort
- ▶ The final loop takes  $O(|E| \cdot t_f + |V| \cdot t_u)$  time

Putting it all together:

$$O(|V| \cdot t_m + |E| \cdot \log(|E|) + |E| \cdot t_f + |V| \cdot t_u)$$

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### The DisjointSet ADT

But wait, what exactly is  $t_m$ ,  $t_f$ , and  $t_u$ ? How exactly do we

implement makeMST(v), findMST(v), and union(u, v)? We can do so using a new ADT called the DisjointSet ADT!

