

# Graphs and Network Flows

## ISE 411

### Lecture 7

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## References for Today's Lecture

- Required reading
  - Chapter 20
- References
  - AMO [Chapter 13](#)
  - CLRS [Chapter 23](#)

# Minimum Spanning Trees

- Optimality Conditions
- Kruskal's Algorithm
- Prim's Algorithm

# Combinatorial Optimization

- A *combinatorial optimization problem* consists of
  - a ground set of elements  $E$ ,
  - an associated set  $\mathcal{F}$  of subsets of  $E$  called the *feasible subsets*.
  - A cost vector  $\mathbb{R}^E$ .
- The cost  $c(S)$  of a feasible subset is  $\sum_{s \in S} c_s$ .
- The goal is to find a subset of minimum cost.

## Minimum Spanning Trees

- Recall that a *spanning tree*  $T$  of  $G$  is a connected acyclic subgraph that spans all the nodes of  $G$ .
- The total cost of a spanning tree is the sum of the costs of the arcs in the tree.
- Given an undirected graph  $G = (N, A)$  with  $n$  nodes and  $m$  arcs and with a length or cost  $c_{ij}$  associated with each arc  $(i, j) \in A$ , the minimum spanning tree problem is to find a spanning tree with the smallest total cost (length).
- This is a combinatorial optimization problem.

## Optimality Conditions

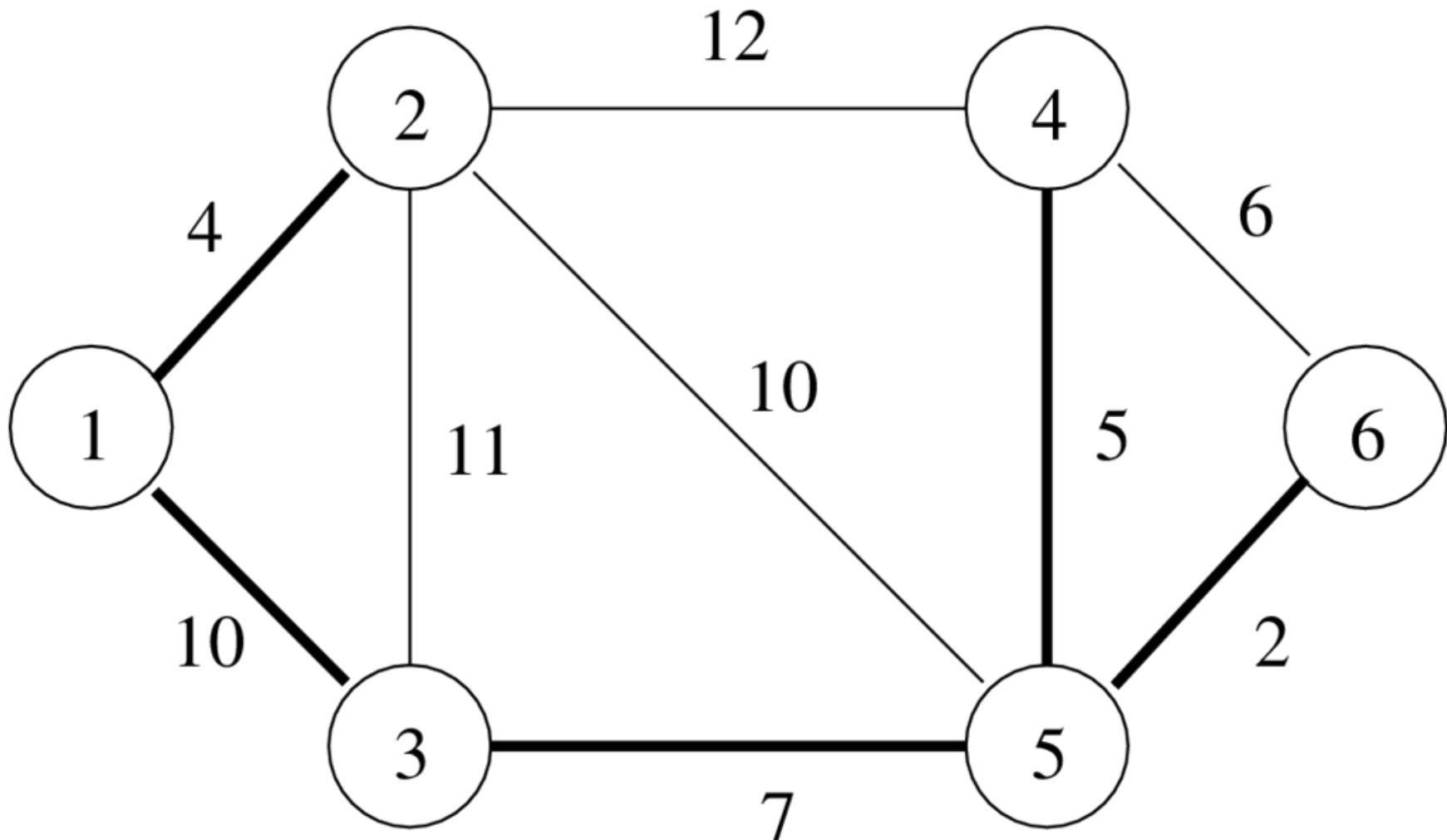
- Cut Optimality Conditions
- Path Optimality Conditions

### Properties of a Spanning Tree

- For every non-tree arc  $(k, l)$ , a spanning tree  $T$  contains a unique path from node  $k$  to node  $l$ . The arc  $(k, l)$  together with the unique path defines a cycle.
- If we delete any tree arc  $(i, j)$  from a spanning tree, we partition the node set into two subsets, which define a cut in the graph.

## Cut Optimality Conditions

**Theorem 1. [13.1]** *A spanning tree  $T^*$  is a minimum spanning tree if and only if for every tree arc  $(i, j) \in T^*$ ,  $c_{ij} \leq c_{kl}$  for every arc  $(k, l)$  contained in the cut formed by deleting arc  $(i, j)$  from  $T^*$ .*



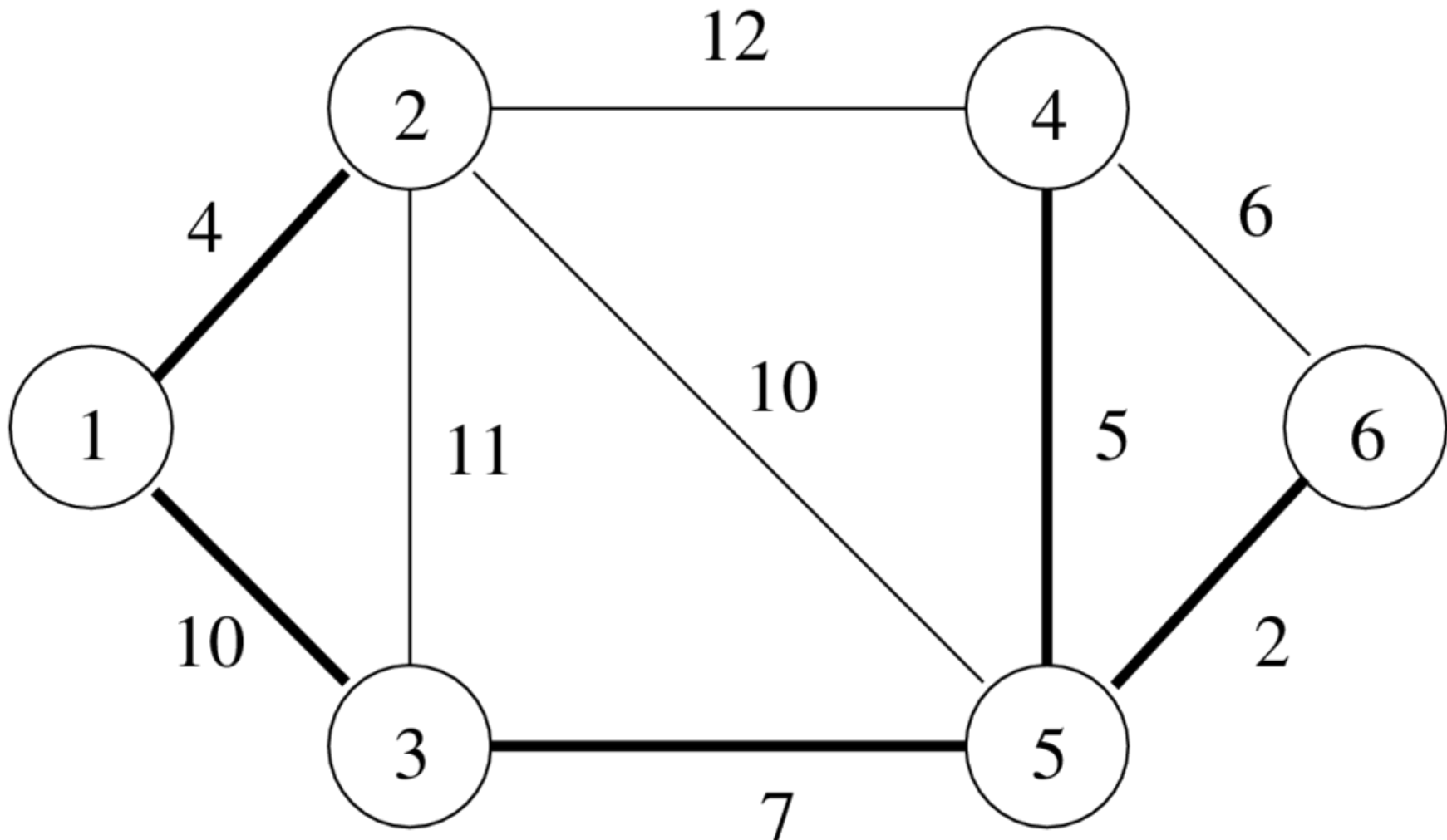
## Proof of Theorem 13.1

1. Show if  $T^*$  is a MST, then  $T^*$  must satisfy the Cut Optimality Conditions.
2. Show if any tree  $T^*$  satisfies the Cut Optimality Conditions, then  $T^*$  is a MST.



## Path Optimality Conditions

**Theorem 2. [13.3]** A spanning tree  $T^*$  is a MST if and only if for every non-tree arc  $(k, l)$  of  $G$ ,  $c_{ij} \leq c_{kl}$  for every arc  $(i, j)$  contained in the path in  $T^*$  connecting nodes  $k$  and  $l$ .



## Proof of Theorem 13.3

1. Show if  $T^*$  is a MST, then  $T^*$  satisfies the Path Optimality Conditions.
2. Show if for every non-tree arc  $(k, l)$  of  $G$   $c_{ij} \leq c_{kl}$  for every arc  $(i, j)$  contained in the path in  $T^*$  connecting nodes  $k$  and  $l$ , then  $T^*$  is a MST.

## Algorithm Based on Cut Optimality

- Prim's algorithm is motivated by the cut optimality conditions.
- We build up the tree one edge at a time as one connected component.
- In each iteration, we will connect one more node to the current tree.
- We do this by adding the edge that is the minimum length edge across the cut induced by the current set of connected nodes.
- Why does this guarantee optimality?
- How do we do this?

## Prim's Algorithm

algorithm *Prim*

$$T = \emptyset$$

$$S = \{1\}; \bar{S} = N - \{1\}$$

while ( $|S| < n$ ) do

    find arc  $(i, j)$  in  $[S, \bar{S}]$  with minimum cost

$$T = T \cup \{(i, j)\}$$

$$S = S \cup \{j\}; \bar{S} = \bar{S} - \{j\}$$

# Complexity

- Number of iterations?
- Dominant step of each iteration?

## Prim's Algorithm

- For each node  $j \in \bar{S}$ 
  - $d(j) = \min$  cost of arcs in the cut incident to a node  $j \notin \bar{S}$
  - $d(j) = \min\{c_{ij} : (i, j) \in [S, \bar{S}]\}$
  - $\text{pred}(j) = i$  such that  $c_{ij} = \min\{c_{ij} : (i, j) \in [S, \bar{S}]\}$ .
- To find min cost arc, compute  $\min\{d(j) : j \in \bar{S}\}$ .
- Suppose  $\hat{j}$  is the min, then  $(\text{pred}(\hat{j}), \hat{j})$  is min cost arc.
- Move  $\hat{j}$  to  $S$  and update distance and predecessor labels for nodes adjacent to  $\hat{j}$ .

## Running Time of Prim's Algorithm

- Note that Prim's Algorithm is a graph search procedure.
- However, the procedure for determining the search order is more complex than previous ones.
- At each step, we need to update some node labels and then be able to determine the node with the minimum label.
- The key to implementing the procedure is an efficient data structure.
- What is the running time for a naive implementation?

## Implementation with Priority Queues

- The running time depends critically on how keep track of the minimum label as the algorithm progresses.
- To get a strongly polynomial time algorithm, we must use a more general data structure for maintaining a *priority queue*.
- For a given order set  $H$ , this data structure should support the operations
  - `push(item, value)` (to add and change value of an item)
  - `peek()`
  - `pop()`



## Binary Heaps

- A *binary heap* is a balanced binary tree with additional structure that allows it to function efficiently as a priority queue.
- The additional structure needed to support these operations is that **each node has a higher priority than either of its children**.
- Balanced binary trees can be stored very efficiently in a single array.
  - The root is stored in position  $0$ .
  - The children of the node in position  $i$  are stored in positions  $2i + 1$  and  $2i + 2$ .
  - This determines a unique storage location for every node in the tree and makes it easy to find a node's parent and children.
  - Using an array, basic operations can be performed very efficiently.

## Creating the Heap

- Any node whose priority is higher than either of its children is said to satisfy the *heap property*.
- Consider a tree in which all nodes except for the root have the heap property.
- We can easily transform this into a tree in which every node has the heap property (*how?*).
- This operation is called *heapify()*.
- By calling *heapify()* on each node, starting at the lowest level and working upward, we can transform an unordered binary tree into a heap.
- This is how we create the initial heap.
- Note that this step is unnecessary for implementing Dijkstra's. Why?

## Operations on a Heap

- The node with the highest priority is always the root.
- To **change the priority** of a node
  
- To **insert** a node
  
- To **delete** a node
  
- What are the running times of these operations?

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# Analyzing Prim's with a Binary Heap

## Algorithm Based on Path Optimality

- Kruskal's algorithm motivated by path optimality conditions.
- We build up the tree one edge at a time, but this time we build multiple components simultaneously.
- In each step, we will add the minimum edge that does not form a cycle with the edges already added.
- Why does this guarantee optimality?
- How do we implement it?

## Kruskal's Algorithm

**algorithm** *Kruskal*

sort edges in non-decreasing order of length

LIST :=  $\emptyset$

while ( $|\text{LIST}| < |N| - 1$  and  $\exists$  unexamined edges) do

$e :=$  unexamined edge with minimum length

    if adding  $e$  to LIST does not create a cycle

        add  $e$  to LIST

    else discard  $e$

## Kruskal's Algorithm: Complexity

- The algorithm has two steps.
  - Sorting the edge list:  $O(m \log m) = O(m \log n)$
  - Building the tree: ??
- To determine which edges we are allowed to add in each step requires a data structure for storing *connected components*.
- The data structure must support two operations.
  - `find(i, j)`: Are *i* and *j* in the same component?
  - `union(i, j)`: Merge the components *i* and *j*.

## Quick Find Implementation of Union-Find

- The simplest implementation involves an array of length  $n$ .
- We will maintain the array such that two items are in the same subset if and only if the array entries are equal.
- This makes the `find(i, j)` constant time, so we call this implementation *quick find*.
- How do we implement `union(i, j)`?
- What is the running time?
- Note that this could also be implemented using linked lists.



## Quick Union Implementation of Union-Find

- To speed up the union operation, we maintain the array in a different fashion.
- We will consider the  $i^{\text{th}}$  entry of the array to be a pointer to another item.
- To perform `find(i, j)`,
  - Follow the pointers from nodes  $i$  and  $j$  until reaching a node that points to itself, called the *representative*
  - If the same representative is reached from both nodes  $i$  and  $j$ , then they are in the same subset.
- To perform `union(i, j)`, perform the find operation and then point the representative for  $i$  to the representative for  $j$ .
- What is the performance now?

## Weighted Quick Union

- Note that the **quick union** algorithm essentially builds a tree out of the nodes in each component, with the root being the representative.
- As in a heap, the running time of the find operation depends on the depth of the trees.
- Each union operation essentially connects two trees together by pointing the root of one tree to the root of the other.
- One way to limit the depth of the tree is to always point the smaller tree to the larger one.
- This ensures that each find takes less than  $\log n$  steps.
- Note that we must now keep track of the number of nodes in each tree, but that's easy to do.
- Another approach is to keep track of the height of each tree and always point the **shorter** tree to the **taller** one.

## Path Compression

- Ideally, we would like each item to point directly to the representative of its subset.
- One possibility is to simply keep track of all the nodes encountered in the path to the root.
- After reaching the root, set all the nodes on the path to point to the root.
- This is easy to implement recursively and doesn't change the asymptotic running time.
- An easier method to implement is *compression by halving*, which is setting each node to point to its grandparent.
- Combining weighted quick union with path compression yields a total running time for connected components of approximately  $O(m)$ .

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# Analyzing Kruskal's with Optimized Union-Find