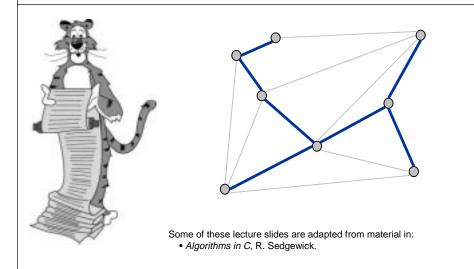
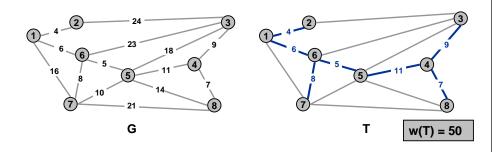
Minimum Spanning Tree

Minimum Spanning Tree



Minimum spanning tree (MST). Given connected graph G with positive edge weights, find a min weight set of edges that connects all of the vertices.



Cayley's Theorem (1889). There are V^{V-2} spanning trees on the complete graph on V vertices.

. Can't solve MST by brute force.

Applications

MST is fundamental problem with diverse applications.

- Designing physical networks.
- telephone, electrical, hydraulic, TV cable, computer, road
 Cluster analysis.
 - delete long edges leaves connected components
 - finding clusters of guasars and Seyfert galaxies
 - analyzing fungal spore spatial patterns
- . Approximate solutions to NP-hard problems.
 - metric TSP, Steiner tree
- Indirect applications.
 - max bottleneck paths
 - describing arrangements of nuclei in skin cells for cancer research
 - learning salient features for real-time face verification
 - modeling locality of particle interactions in turbulent fluid flow
 - reducing data storage in sequencing amino acids in a protein

Optimal Message Passing

Optimal message passing.

- Distribute message to N agents.
- Each agent can communicate with some of the other agents, but their communication is (independently) detected with probability p_{ii}.
- Group leader wants to transmit message to all agents so as to minimize the total probability that message is detected.

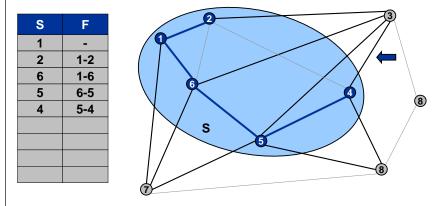
Objective.

- Find tree T that minimizes: $1 \prod_{(i,j) \in T} (1 p_{ij})$
- Or equivalently, that maximizes: $\prod_{(i,j)\in T} (1-p_{ij})$
- Or equivalently, that maximizes: $\sum_{(i,j)\in T} \log(1-p_{ij})$
 - MST with weights = log (1 p_{ii}) weights p_{ii} also work!

Prim's Algorithm

Prim's algorithm. (Jarník 1930, Dijkstra 1957, Prim 1959)

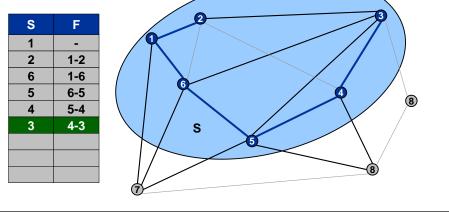
- . Initialize F = $\varphi,\,$ S = {s} for some arbitrary vertex s.
- Repeat until S has V vertices:
 - let f be smallest edge with exactly one endpoint in S
 - add other endpoint to S
 - add edge f to F



Prim's algorithm. (Jarník 1930, Dijkstra 1957, Prim 1959)

Prim's Algorithm

- Initialize $F = \phi$, $S = \{s\}$ for some arbitrary vertex s.
- Repeat until S has V vertices:
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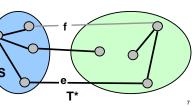
Prim's Algorithm: Proof of Correctness

Theorem. Upon termination of Prim's algorithm, F is a MST.

Proof. (by induction on number of iterations)

Invariant: There exists a MST T* containing all of the edges in F.

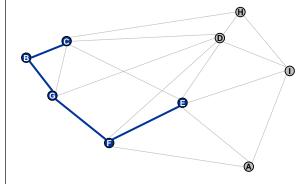
- Base case: $F = \phi \implies$ every MST satisfies invariant.
- Induction step: true at beginning of iteration i.
 - at beginning of iteration i, let S be vertex subset and let f be the edge that Prim's algorithm chooses
 - if $f \in T^*$, T^* still satisfies invariant
 - o/w, consider cycle C formed by adding f to T*
 - let $\textbf{e} \in \textbf{C}$ be another arc with exactly one endpoint in S
 - $\mathbf{c}_{f} \leq \mathbf{c}_{e}~$ since algorithm chooses f instead of e
 - e ∉ F by definition of S
 - T* \cup { f } { e } satisfies invariant



Prim's Algorithm: Classic Implementation

Use adjacency matrix.

- . S = set of vertices in current tree.
- For each vertex not in S, maintain vertex in S to which it is closest.
- . Choose next vertex to add to S using min dist[w].
- Just before adding new vertex v to S:
 - for each neighbor ${\bf w}$ of ${\bf v},$ if ${\bf w}$ is closer to ${\bf v}$ than to a vertex in S, update dist[w]

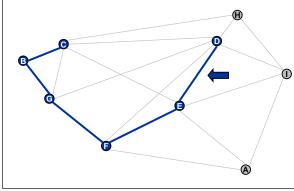




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Vertex	Nearest	Dist	
Α	Е	15	
В	-	-	
С	-	-	
D			
E	-	-	
F	-	-	
G	-	-	
Н	D	4	
Ι	D	6	

Prim's Algorithm: Classic Implementation

Running time.

• V - 1 iterations since each iteration adds 1 vertex.

Each iteration consists of:

- . Choose next vertex to add to S by minimum dist[w] value.
 - O(V) time.
- For each neighbor w of v, if w is closer to v than to a vertex in S, update dist[w].

- O(V) time.

O(V²) overall.

Prim's Algorithm: Priority Queue Implementation

Prim's Algorithm pseudocode				
$Q \leftarrow PQinit()$ for each vertex v in graph G key(v) $\leftarrow \infty$				
pred(v) ← nil PQinsert(v, Q)				
<pre>key(s) ← 0 while (!PQisempty(Q)) v = PQdelmin(Q) for each edge v-w s.t. w is in Q if key(w) > c(v,w) PQdeckey(w, c(v,w), Q) pred(w) ← v</pre>				

Prim's Algorithm: Priority Queue Implementation

Analysis of Prim's algorithm.

- PQinsert(): V vertices.
- PQisempty(): V vertices.
- PQdelmin(): V vertices.
- PQdeckey(): E edges.

	Priority Queues		
Operation	Array	Binary heap	Fibonacci heap*
insert	N	log N	1
delete-min	N	log N	log N
decrease-key	1	log N	1
is-empty	1	1	1
Prim	V ²	E log V	E + V log V

PFS vs. Classic Prim

Which algorithm is faster?

- . Classic Prim: O(V²).
- Prim with binary heap: O(E log V).

Answer depends on whether graph is SPARSE or DENSE.

- 2,000 vertices, 1 million edges
 - Heap: 2-3 times SLOWER
- 100,000 vertices, 1 million edges
 Heap: 500 times FASTER
- 1 million vertices, 2 million edges
 - Heap: 10,000 times FASTER.

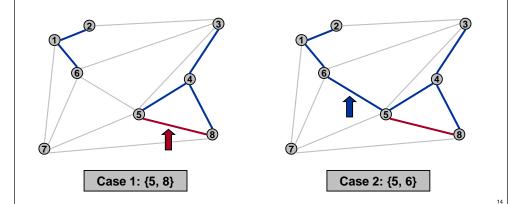
Bottom line.

- . Classic Prim is optimal for dense graphs.
- . Heap implementation far better for sparse graphs.

Kruskal's Algorithm

Kruskal's algorithm (1956).

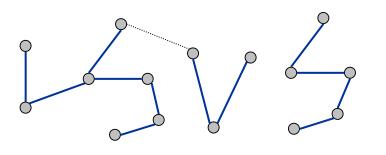
- . Initialize $F = \phi$.
- Consider arcs in ascending order of weight.
- If adding edge to forest F does not create a cycle, then add it.
 Otherwise, discard it.



Kruskal's Algorithm: Implementation

How to check if adding an edge to F would create a cycle?

- Naïve solution: use depth first search.
- Clever solution: use union-find data structure from Lecture 1.
 - each tree in forest corresponds to a set
 - to see if adding edge between ${\bf v}$ and ${\bf w}$ creates a cycle, check if ${\bf v}$ and ${\bf w}$ are already in same component
 - when adding v-w to forest F, merge sets containing v and w



Kruskal's Algorithm: C Implementation

Kruskal's Algorithm

```
// Fill up mst[] with list of edges in MST of graph G
void GRAPHmstE(Graph G, Edge mst[]) {
   int i, k, v, w;
   Edge a[MAXE];
                               // list of all edges in G
   int E = GRAPHedges(a, G); // # edges in G
   sort(a, 0, E-1);
                              // sort edges by weight
  UFinit(G->V);
   for (i = k = 0; i < E \&\& k < G \rightarrow V-1; i++)
      v = a[i].v;
      w = a[i].w;
      // if edge a[i] doesn't create a cycle, add to tree
      if (!UFfind(v, w)) {
         UFunion(v, w);
         mst[k++] = a[i];
   }
```

15

Kruskal's Algorithm: Proof of Correctness

Theorem. Upon termination of Kruskal's algorithm, F is a MST.

Proof. Identical to proof of correctness for Prim's algorithm except that you let S be the set of nodes in component of F containing v.

Corollary. "Greed is good. Greed is right. Greed works. Greed clarifies, cuts through, and captures the essence of the evolutionary spirit."



Gordon Gecko (Michael Douglas)



Kruskal's Algorithm: Running Time

Kruskal analysis. O(E log V) time.

- $O(E \log E) = O(E \log V).$ Sort():
- . UFinit(): V singleton sets.
- . UFfind(): at most once per edge.
- UFunion(): exactly V 1 times.

If edges already sorted. O(E log* V) time.

- . Any sequence of M union-find operations on N elements takes O(M log* N) time.
- In this universe, $\log^* N \le 6$.

Advanced MST Algorithms

Prim, Kruskal, Boruvka.

Fredman-Tarjan (1987).

Deterministic comparison based algorithms.

- O(E log V)
- O(E log log V).
- O(E log*V).
- O(E log (log*V)).
- O(E α (E, V)).
- O(E).

- Chazelle (2000).



Cheriton-Tarjan (1976), Yao (1975).

Gabow-Galil-Spencer-Tarjan (1986).



Worth noting.

- O(E) randomized.
- O(E) verification.
- Karger-Klein-Tarjan (1995). Dixon-Rauch-Tarjan (1992).