

## Kruskals Algorithm

# Minimum Spanning Trees (MST), Kruskal, Prim & Disjoint Sets — Exam Notes

## What is Kruskal's algorithm used for?

Task: Build a Minimum Spanning Tree (MST) of a connected, weighted, undirected graph — i.e., connect all vertices with V-1 edges, no cycles, minimum total weight. lecture13\_kruskals

## Kruskal's process (high level)

- 1. Sort all edges by increasing weight.
- 2. Start with a **forest** of single-vertex trees (each vertex alone).
- Scan edges in order; add an edge iff it connects two different trees (doesn't form a cycle).
- 4. Stop when you've added **V-1** edges → that set is an MST. lecture13\_kruskals

#### **Correctness (simple English):**

At the moment Kruskal considers an edge that connects two components, that edge is the **lightest** across the "cut" separating those components; by the **cut property**, it's safe (belongs to some MST). Replacing any heavier cross-cut edge in a supposed MST with this lighter one can only improve or match the weight, so Kruskal never makes a bad choice.

lecture13\_kruskals

### Prim's algorithm — what, how, why, performance

What: Another MST algorithm that grows a single tree instead of a forest.

#### How (binary heap version):

- Start from any vertex with an empty tree.
- Maintain a priority queue (min-key) of cut edges that connect the current tree to outside vertices.
- Repeatedly extract the minimum-weight eligible edge and add that new vertex and its edges to the queue.
- When all vertices are added → MST, lecture13\_kruskals

#### Why correct (plain English):

At each step, Prim picks the **lightest edge leaving the current tree**. By the **cut property** (the tree vs. the rest is a cut), that lightest leaving edge is always safe to add; thus every step preserves optimality until the whole graph is spanned.

#### Performance (binary heap / PQ):

- Using a binary heap: O(|V| log |V| + |E| log |V|). (Insert/update |E| edges; |V| extractions.) lecture13\_kruskals
- (For context you can remember: an adjacency-matrix + no-heap variant is O(V²), but the heap version above is what you'll usually cite.)

## Kruskal — performance & space

- With efficient DSU: sort edges O(E log E); each find/union is ~amortized
   O(α(V)) → dominated by sorting ⇒ O(E log E) overall. Space O(V + E).
   lecture13\_kruskals
- Naïve implementation (why it's bad):
  - Store a leader[vertex] array and, on union, scan all vertices to relabel (O(V) per successful union).
  - o With  $\sim$ V−1 unions this costs  $O(V^2)$ ; plus sorting  $O(E \log E) \rightarrow O(V^2 + E \log E)$  overall. Space still O(V + E). lecture13\_kruskals

## Fixing the naïve approach: Disjoint Sets (Union-Find)

#### The operations & what we want

- find(x): returns the representative (leader) of x's set → lets us test "same component?".
- union(x, y): merges the two sets containing x and y.

Kruskal needs these operations to be very fast. lecture13\_kruskals

#### Forest representation (parent pointers)

- Maintain a forest where each set is a tree; every node stores a parent; roots are leaders (parent = self).
- union(A,B): make the root of one tree point to the root of the other → O(1)
  pointer change.
- find(x): walk parent pointers to the root → O(height) of the tree (can be O(N) in the worst case if trees get tall). lecture13\_kruskals

#### Union by height (a.k.a. union by rank) — why this gives log V

**Idea:** Keep trees **shallow** by always attaching the **shorter** tree under the **taller** tree.

- A tree's height increases by 1 only when you merge two trees of equal height.
- Induction/doubling argument: the minimum number of nodes in a tree of height h is 2<sup>h</sup> (because to get height h you had to combine two height h-1 trees, each with at least 2<sup>h</sup>(h-1) nodes).
- Therefore with n total nodes, the height can be at most [log<sub>2</sub> n].
- That makes find worst-case O(log V) (and union is one or two finds plus a constant). lecture13\_kruskals

#### Path compression — how it speeds things up

**Mechanism:** After you do find(x) and discover the root r, rewrite every node on the path from x to r so its parent points directly to r.

• This **flattens** the tree along that path.

- A single find might still take up to O(log V), but it makes **future finds** on those nodes **O(1)** until another union changes the structure.
- Combining union by rank (height) + path compression yields a structure where any sequence of m operations on n elements costs O(m · α(n)), where α is the inverse Ackermann function (grows slower than log\*; ≤ 4 for all practical n). In practice this is near-constant time per op. lecture13\_kruskals

Why "amortized" and not worst-case per op?

A costly find happens rarely and **pays for itself** by flattening paths, so the average over many ops becomes tiny. The formal proof (CLRS §19.4) is beyond scope, but the slides give the result and intuition. lecture13 kruskals

#### Bottom line for Kruskal with DSU

Sort edges O(E log E); each of the E find/union ops is ~O(α(V)) amortized ⇒ total O(E log E). This is the standard bound you quote. lecture13\_kruskals

### Proof ideas you can write fast in an exam

#### Kruskal (cut property, plain English)

When Kruskal picks an edge connecting two components, it's the cheapest
edge across that cut; any MST must use some crossing edge, and if it used a
heavier one, swapping in Kruskal's lighter edge would not hurt and can
improve. Thus every chosen edge is safe, and after V-1 such choices we have
an MST. lecture13\_kruskals

#### Prim (same cut property)

 The current tree vs. the rest defines a cut. The lightest outgoing edge is safe by the cut property; adding it preserves optimality until all vertices are included. (Prim recap and complexity on slides.) lecture13\_kruskals

# Pseudocode sketches (what data structures are doing) Kruskal (with DSU):

```
sort edges by weight
make-set(v) for all vertices v
mst_weight = 0
for (u,v,w) in edges:
   if find(u) != find(v): # different components?
      mst_weight += w
      union(u, v)
return mst_weight
```

- Data structures: array/list of edges; DSU with parent[], rank[], path compression in find.
- Flow: sort → scan → conditionally union. lecture13\_kruskals

#### Prim (binary heap):

```
pick any start vertex s

dist[v] = +∞ for all v; dist[s] = 0

push (0, s) into min-heap

while heap not empty:

(key,u) = extract-min()

if u already in tree: continue

add u to tree; add key to total

for each edge (u,v,w):

if v not in tree and w < dist[v]:

dist[v] = w

push (w, v) into heap
```

- Data structures: adjacency list; min-heap keyed by best edge weight to connect each outside vertex.
- Flow: grow one tree, always add the cheapest connecting edge. lecture13\_kruskals

## Naïve vs. optimized — time & space summary

Algorithm / DS	Time	Space	Notes
Kruskal (naïve "leader relabel" union)	O(V <sup>2</sup> + E log E)	O(V+E)	Union relabels whole set each time (O(V)). lecture13_kruskals
Kruskal + DSU (rank + path compression)	O(E log E) overall (sorting dominates)	O(V+E)	Each find/union amortized <b>O(α(V))</b> . lecture13_kruskals
Prim (binary heap)	**O(	V	log

## Quick "why" bullets for DSU complexities (write these if asked "explain why"):

- Union by height/rank ⇒ O(log V) find (worst-case): Height increases only when merging equal-height trees; minimal size doubles each increase ⇒ height ≤ log<sub>2</sub> n. lecture13\_kruskals
- Path compression  $\Rightarrow$  amortized near-O(1): Each expensive find flattens its path; future find s are cheap. Over m operations the total time is O(m  $\alpha$ (n)), with  $\alpha \le 4$  in practice. lecture13\_kruskals