Algorithmics and Programming III

FIB

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- 1 Motivation
 - Example: Coin Exchange
 - What are Greedy Algorithms?
- 2 Scheduling
 - Interval Scheduling Problem
 - Interval Partitioning Problem
 - Lateness Minimization Problem
- 3 Dijkstra's Algorithm
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 - Proof of Dijkstra's Algorithm
- 4 Minimum Spanning Trees
 - General Algorithm
 - Prim's Algorithm
 - Kruskal's Algorithm

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- What if we add a coin of 8 cents?
 - The previous greedy algorithm uses 4 coins: 50 + 10 + 2 + 1
 - However, there is a solution with only 3 coins: 50 + 8 + 5

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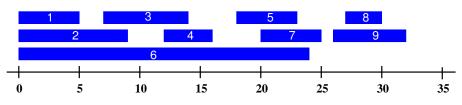
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- In the following we will:
 - Introduce and revisit greedy algorithms for well-known problems
 - Prove them correct using a variety of techniques

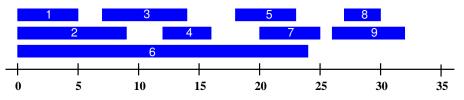
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- Imagine we have a single lab room and N courses that request to use it
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- Requests of two overlapping courses cannot be both met
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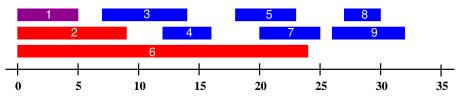
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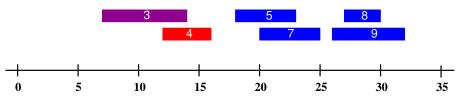
 \bullet {1,3,5,8} or {2,4,7,9} are feasible choices. Are they optimal?



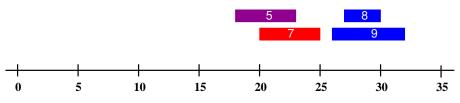
- Idea: start using the room as soon as possible
- Algorithm: while there are available courses, select the available course i that starts as soon as possible. If there is a tie, pick the one with the lowest index. Mark all courses that overlap with i as non-available.
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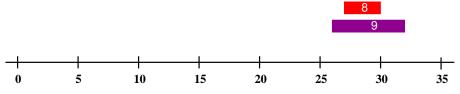
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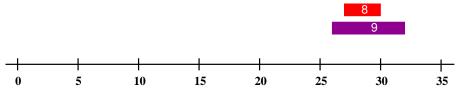
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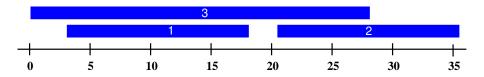
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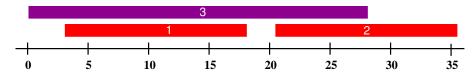
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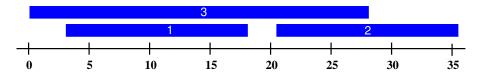
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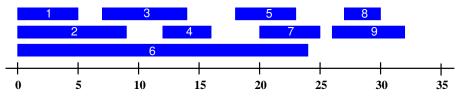
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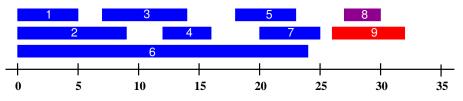
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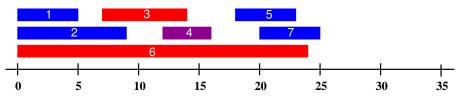
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- But {1,2} is a better solution. So this greedy algorithm is not correct.



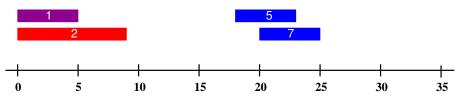
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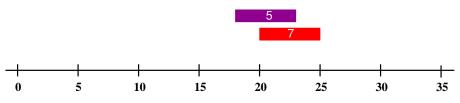
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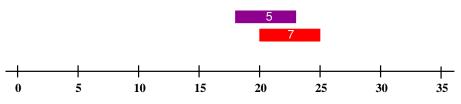
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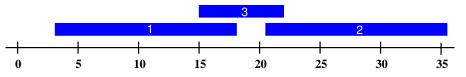
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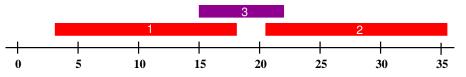
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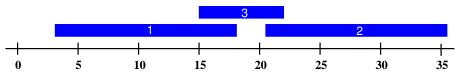
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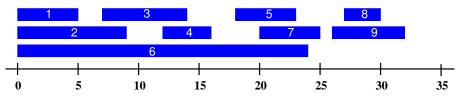
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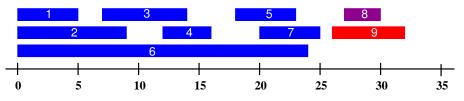
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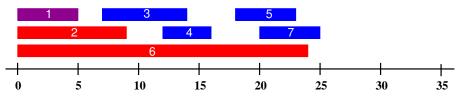
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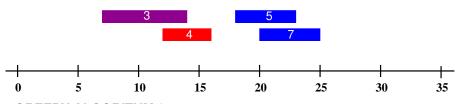
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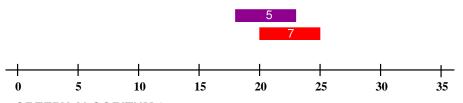
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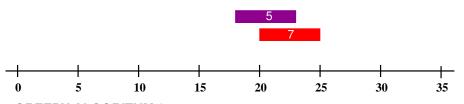
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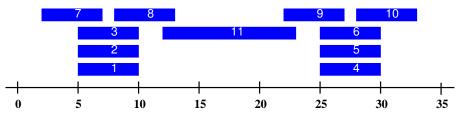
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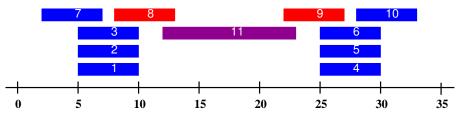
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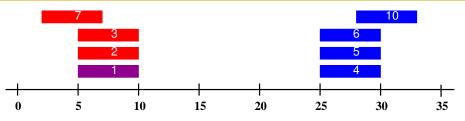
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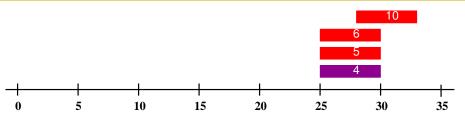
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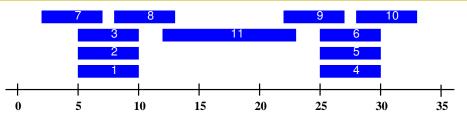
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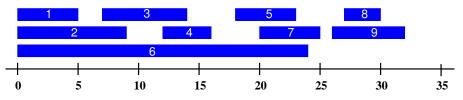
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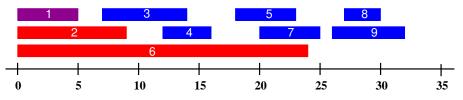
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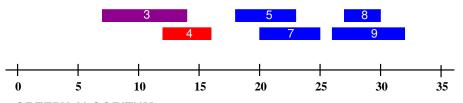
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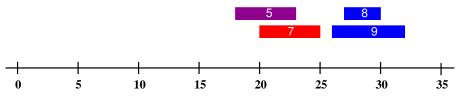
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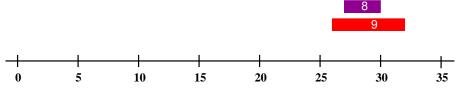
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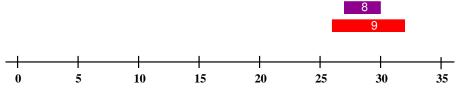
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We will prove that the previous algorithm is correct. Our strategy is the following:

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- We know that $k \le m$ because \mathcal{O} is optimal.

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- We know that $k \le m$ because \mathcal{O} is optimal.
- To see $k \ge m$ we will prove that our algorithm always "stays ahead": for all $1 \le r \le k$ we have $f(a_r) \le f(o_r)$.

- Let S be an arbitrary input set of courses, for which the previous algorithm returns a subset of compatible courses $A = \{a_1, a_2, \dots, a_k\}$.
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- The intuition is that at each step our algorithm has a better subproblem, as it works with courses in the interval $[f(a_r), \ldots]$ rather than in $[f(o_r), \ldots]$

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So when our algorithm selected a_r , one of the candidates to choose was o_r . As it chooses the one with the smallest finishing time, we have $f(a_r) \le f(o_r)$.

Remember:

$$A = \{a_1, a_2, \dots, a_k\}$$
 (the algorithm solution)

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 (optimal)

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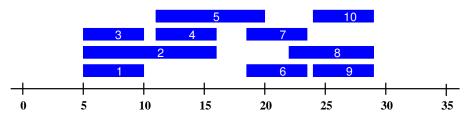
As $f(a_1) \leq \ldots \leq f(a_k)$, we have o_{k+1} does not overlap with any of a_1, \ldots, a_k . So o_{k+1} was still available when our algorithm finished: contradiction! (because it finishes when no available course exists).

```
struct Task {
  int id;
  int starting;
  int finishing;
};
bool before (const Task& t1, const Task& t2) {
  return t1.finishing < t2.finishing;
int main() {
  vector<Task> tasks;
  int s, f, id = 1;
  while (cin >> s >> f) {
    tasks.push_back(Task{id,s,f});
    ++id;
  sort(tasks.begin(), tasks.end(), before);
  vector<int> result = interval_scheduling(tasks);
  for (auto& x: result) cout << x << endl;
```

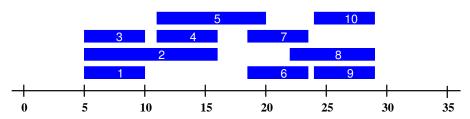
```
// PRE: tasks are sorted increasingly by finishing time
vector<int> interval_scheduling(const vector<Task>& tasks) {
  int n = tasks.size();
 int idx = 0:
  int last_f = INT_MIN; // Acts as -\infty
  while (idx < n) {
   // Look for non-overlapping task w/ smallest finishing
   while (tasks[idx].starting < last_f and idx < n) ++idx;
    if (idx < n) { // Add tasks[idx] to solution
     res.push_back(tasks[idx].id);
     last_f = tasks[idx].finishing;
  return res;
```

- Imagine we have N courses that need to be allocated to some rooms
- Again, we know starting time s(i) and finishing time f(i) of each course
- The difference is that now:
 - we have to allocate all courses
 - we have an unlimited number of rooms
- We wish to find out the minimum number of rooms needed so that no two overlapping courses are assigned to the same room

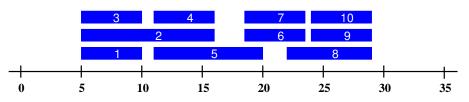
Scheduling using 4 rooms:



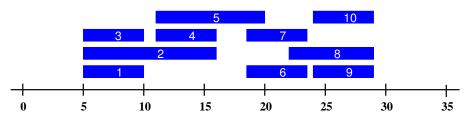
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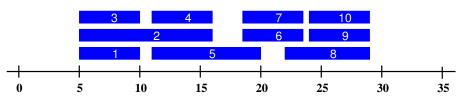
Scheduling using 3 rooms:



Scheduling using 4 rooms:



Scheduling using 3 rooms:



 \bullet Latter is optimal, as $\{1,2,3\}$ are overlapping and hence need 3 rooms.

- Let us define the depth of the problem as the maximum number of courses that overlap on any instant of time
- **CLAIM:** The number of rooms needed is ≥ the depth of the problem
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- We should prove:
 - In the second step there is always at least one free room
 - No overlapping courses get the same rooms (easy!)

Proposition

In the second step of the algorithm there is always one free room to choose

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Assume it is not. Let c be the first course that has no free room to choose.

This means that at time s(c),

all d rooms are occupied by previous courses that are active at that moment.

Those courses, together with c, form a set of d+1 overlapping courses, which is a contradiction with the definition of the depth d.

```
bool overlap(const Task& t1, const Task& t2) {
  return t1.finishing > t2.starting and
         t2.finishing > t1.starting; }
bool before (const Task& t1, const Task& t2) {
  return t1.starting < t2.starting; }
int main() {
 vector<Task> tasks;
  int s, f, id = 1;
  while (cin >> s >> f) {
    tasks.push_back(Task{id,s,f});
    ++id;
  sort(tasks.begin(),tasks.end(), before);
  vector<int> labelling = interval_part(tasks);
  for (uint i = 0; i < tasks.size(); ++i)
  cout << "Task " << tasks[i].id << " gets room " <<
     labelling[i] << endl;</pre>
```

```
// PRE: tasks are sorted increasingly by starting time
vector<int> interval_part (const vector<Task>& tasks) {
  int n = tasks.size();
  vector<int> labelling(n,-1); // -1 for unassigned label
  for (int i = 0; i < n; ++i) {
    // Mark not available labels for the i-th task
    vector<int> available_label(n, true); // large UB
    for (int j = 0; j < i; ++j)
      if (overlap(tasks[i], tasks[j]))
         available_label[labelling[j]] = false;
    int 1 = 0:
    while (not available label[1]) ++1;
    labelling[i] = 1;
  return labelling;
```

- Imagine we have a set of N tasks that request to use the same resource, of which we only have one
- For each task i we know its duration du(i) and its deadline d(i) (ideally, the task should be finished before this time)

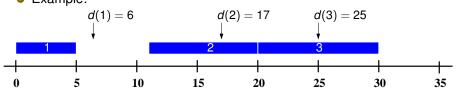
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- Our goal is to find starting times for all tasks so as to minimize the maximum lateness, i.e.

 $\min \max_{i=1}^{N} \ell(i)$

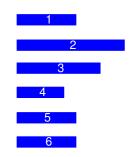
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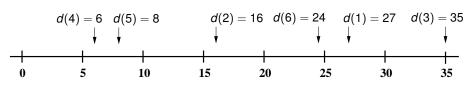
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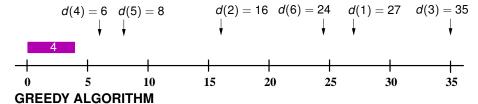
Example:



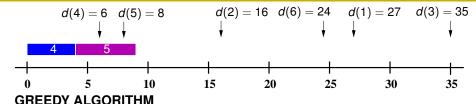
This non-optimal schedule has a maximum lateness of 5



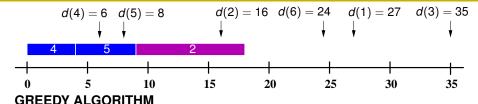




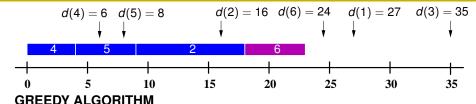
- Idea: schedule first the tasks with earliest deadline
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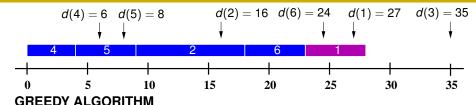
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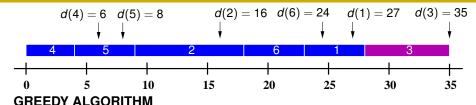
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 - Ohoose task 2. Starts at 9, finishes at 18. Lateness is 2.



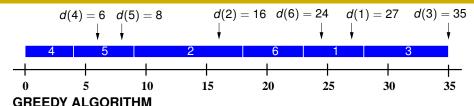
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 - Choose task 1. Starts at 23, finishes at 28. Lateness is 1.



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 - Ohoose task 1. Starts at 23, finishes at 28. Lateness is 1.
 - Ohoose task 3. Starts at 28, finishes at 35. Lateness is 0. Maximum lateness is 2

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- 2 We prove there's an optimal schedule with no inversions and no idle time

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All schedules with no inversions and no idle time have the same max lateness

Proof: In a schedule with no inversions, if d(i) > d(j) then s(i) > s(j), i.e., task i necessarily follows task j.

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In both schedules tasks with the same deadline d will appear consecutively (after all tasks with deadlines < d and before all tasks with deadlines > d), but maybe in a different order.

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This happens for all deadlines d, so the two schedules have the same maximum lateness.

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If it has no inversions, we are done.

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- will have one less inversion
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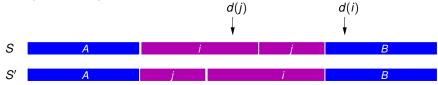
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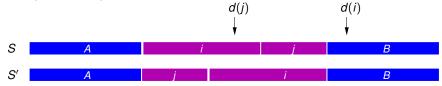
By repeating this swapping process, we will end up having an optimal schedule with no inversions.

Proof (continued):



S' has one less inversion than S:

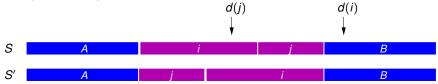
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S' has one less inversion than S:

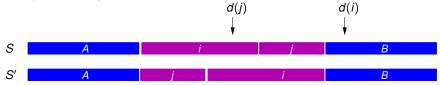
No new inversions among tasks in A or B are created. Inversions in S' of the form (k, j), (k, i) with k in A belong to S too. Inversions in S' of the form (i, k), (j, k) with k in B belong to S too.

Proof (continued):



• The maximum lateness of S' is no larger than the one of S:

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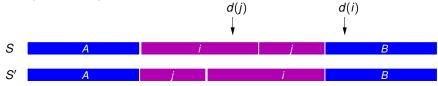


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In S' only the finishing times of i and j are changed.

As the finishing time of *j* is smaller, only *i* may have a larger lateness

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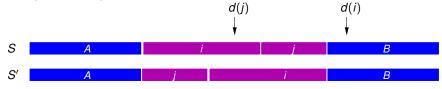
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If *i* is late in S', its lateness is s'(i) + du(i) - d(i) = s(j) + du(j) - d(i).

But since d(i) > d(j), we have i cannot be later in S' than j in S:

$$\ell'(i) = s(j) + du(j) - d(i) < s(j) + du(j) - d(j) = \ell(j).$$

```
struct Task {
  int id;
  int duration:
  int deadline;
  Task(int i, int du, int de):
  id(i), duration(du), deadline(de){} };
bool before (const Task& t1, const Task& t2) {
  return t1.deadline < t2.deadline; }
int main() {
  vector<Task> tasks;
  int du, de, id = 1;
  while (cin >> du >> de) {
    tasks.push_back(Task(id,du,de)); ++id;}
  sort(tasks.begin(), tasks.end(), before);
  vector<int> starting_time = lateness(tasks);
  for (uint i = 0; i < tasks.size(); ++i)
     cout << "Task " << tasks[i].id << " starts at " <<
         starting_time[i] << endl;
```

```
// PRE: tasks is sorted increasingly by deadline
vector<int> lateness(const_vector<Task>& tasks) {
  int n = tasks.size();
  vector<int> starting_time(n);
  int t = 0;
  for (int i = 0; i < n; ++i) {
    starting_time[i] = t;
    t += tasks[i].duration;
  return starting time;
```

Chapter 4. Greedy Algorithms

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Dijkstra's Algorithm

Dijkstra's algorithm for shortest paths

```
Input: Graph G = (V, E), directed or not;
       non-negative weights in the edges \{w(u, v) \mid (u, v) \in E\};
       initial vertex s:
Output: For all vertex u reachable from s,
       dist(u) = distance from s to u (denoted in the following by \delta(s, u))
DIJKSTRA (G, w, s)
   for all vertex u \in V do dist(u) = \infty
   dist(s) = 0
   H = \text{create-priority-queue}(V)
                                                 (using dist as keys)
   while not empty(H)
       u = \text{remove-min}(H)
                                          ← This is the greedy part
      for all edge (u, v) \in E do
          if dist(v) > dist(u) + w(u, v)
                                                         (relax edge)
              dist(v) = dist(u) + w(u, v)
             decrement-key(H, v)
```

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If $\mathit{dist}(u){<}\infty$, then $\mathit{dist}(u)$ corresponds to the length of some path from s to u

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Consider now the *k*-th update.

Assume the property holds up to the k-1-th update.

The *k*-th update is done with the command

$$dist(v) = dist(u) + w(u, v)$$

Since $dist(u) < +\infty$ corresponds to the length of some path from s to u, and there is an edge $u \to v$ with weight w(u, v), the property is true.

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Corollary

For all vertices u, it always holds that $dist(u) \ge \delta(s, u)$.

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We also know that $u \neq s$.

Now let us consider the shortest path from s to u.

Before we removed u from H, this path went from $s \notin H$ to $u \in H$.

Let $x \to y$ be the first edge in this path from a vertex not in H to a vertex in H. (Note that it could be s = x or y = u).

This path is of the form $s \rightsquigarrow x \rightarrow y \rightsquigarrow u$.

Proof (continued):

Let us consider the state of the algorithm right after x was removed from H. When processing edge $x \to y$, two things may happen:

- If $dist(y) \leq dist(x) + w(x, y)$, nothing is done
- If dist(y) > dist(x) + w(x, y), the edge is relaxed and dist is updated so that dist(y) = dist(x) + w(x, y).

So after processing edge (x, y) we have $dist(y) \le dist(x) + w(x, y)$.

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This property is preserved from that moment on:

- dist(x) is not changed $(dist(x) = \delta(s, x))$, so it cannot decrease)
- dist(y) can only decrease

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Since $s \rightsquigarrow x \to y$ is part of the shortest path between s and u, it is the shortest path between s and y and so $\delta(s,y) = \delta(s,x) + w(x,y)$.

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from which we conclude dist(y) < dist(u) just before u was removed from H.

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Dijkstra's Algorithm computes the distances from s to all reachable vertices

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But at the end of the algorithm, *H* is empty.

As all vertices are eventually removed, all distances are duly computed

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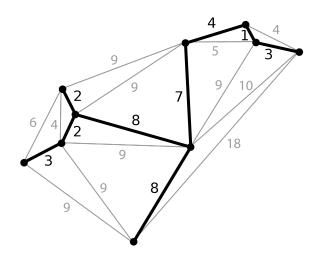
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Definition

A minimum spanning tree (MST) of G is a spanning tree A of G such that its total weight

$$\omega(A) = \sum_{e \in A} \omega(e)$$

is minimum among all spanning trees of G



Source: https://commons.wikimedia.org/wiki/File:Minimum_spanning_tree.svg

There are many different algorithms for computing MST's.

They typically follow the scheme:

```
A = \emptyset; Cand = E; while (|A| \neq |V| - 1) { choose e \in Cand that does not close a cycle in A; A = A \cup \{e\}; Cand = Cand - \{e\}}
```

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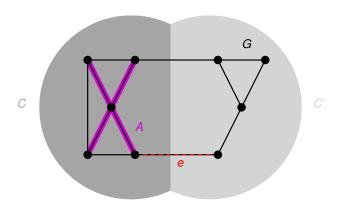
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An edge e respects a cut (C, C') if the ends of e belong both to C or to C'; otherwise, we say that e crosses the cut.

Theorem

Let *A* be a promising set of edges that respects a cut (C, C') of *G*. Let *e* be an edge of minimum weight among those that cross the cut (C, C'). Then $A \cup \{e\}$ is promising.



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This theorem gives a recipe for designing algorithms for MST:

- start with an empty set of edges A
- define a cut of G which is respected by A
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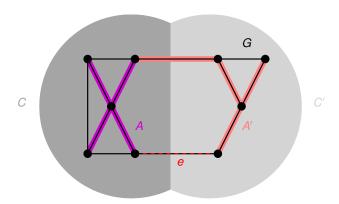
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Any algorithm that follows this scheme is automatically correct.

Note that step 3 chooses the edge greedily (the one with minimum weight among those that cross the cut)

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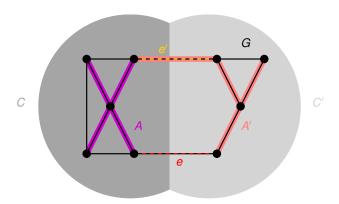
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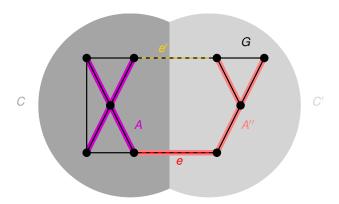
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As A respects the cut, there is an edge $e' \in A' - A$ that crosses the cut (otherwise, A' would not be connected).

The subgraph $A'' = A' - \{e'\} \cup \{e\}$ is a spanning tree and

$$\omega(A') \le \omega(A'') = \omega(A') - \omega(e') + \omega(e)$$

Hence $\omega(e') \leq \omega(e)$.

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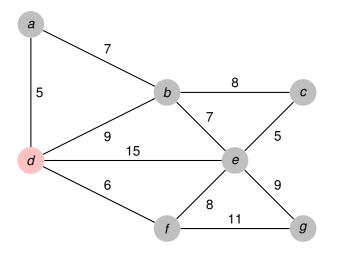
Since A'' is a MST, $A \cup \{e\}$ is promising.

In Prim's Algorithm (also known as Prim-Jarník's Algorithm), we keep a subset of visited vertices.

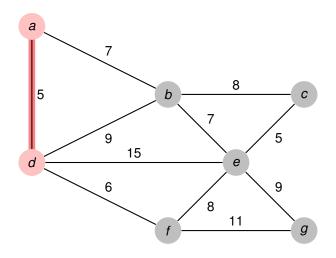
The set of vertices is thus partitioned into visited and non-visited vertices.

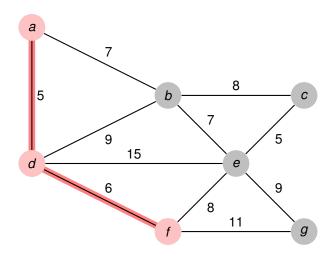
Each iteration of the algorithm chooses an edge of minimum weight among those that join a visited vertex and a non-visited one.

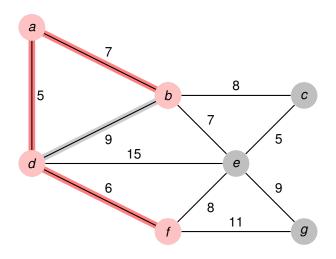
By the theorem, the algorithm is correct.

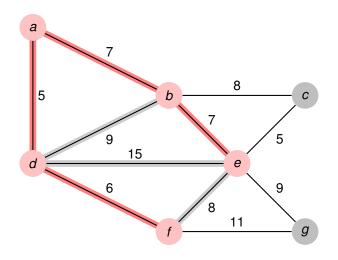


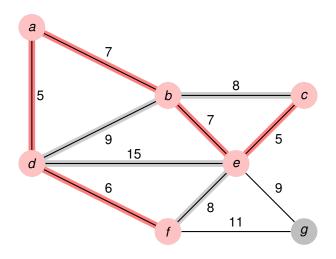
 $\textbf{Font:} \ \texttt{www.texample.net/tikz/examples/author/kjell-magne-fauske}$

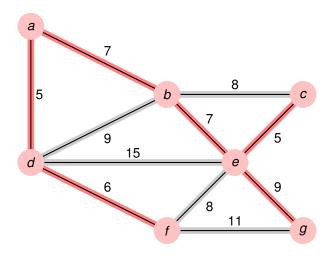












```
typedef pair < double, pair <int, int > WEdge;
typedef vector< vector< pair<double, int> > > WGraph;
int Prim(const WGraph& G, vector<pair<int,int>>& MST) {
  vector<bool> used(G.size(), false);
  priority_queue<WEdge, vector<WEdge>, greater<WEdge>> Q;
  used[0] = true;
  for (auto& x : G[0]) Q.push({x.first, {0, x.second}});
  int totalWeight = 0; uint sz = 1;
  while (sz < G.size()) {
    int u = Q.top().second.first;
    int v = Q.top().second.second;
    double wEdge = Q.top().first; Q.pop();
    if (not used[v]) {
      used[v] = true;
      MST.push_back({u,v});
      totalWeight += wEdge; ++sz;
      for (auto e : G[v]) Q.push({e.first, {v, e.second}});
  return totalWeight; }
```

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Prim's Algorithm runs in $O(m \log n)$ time

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In the implementation, the cost of the loop dominates overall cost.

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There are O(m) iterations.

The choice of the edge at each iteration takes time $O(\log m)$.

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Prim's Algorithm runs in $O(m \log n)$ time

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In the implementation, the cost of the loop dominates overall cost.

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In total:

$$O(m\log m) + \sum_{x \in V} O(\deg(x) \log m) = O(m\log m) = O(m\log n)$$

Kruskal's Algorithm

In Kruskal's Algorithm, at each step we keep:

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Next we will see the algorithm is correct, as it follows the previous recipe:

- start with an empty set of edges A
- define a cut of G which is respected by A
- 3 choose the edge e with minimum weight among those that cross the cut
- add e to A

We only need to determine what is the cut of Kruskal's algorithm

Let G = (V, E) be the graph for which we want to find an MST.

At each step, consider $\{u, v\}$ the edge added by Kruskal (i.e. the edge in P of minimum weight that does not create a cycle in SF)

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We define the cut as follows:

$$C = \{ w \in V \mid u \text{ has a path to } w \text{ in } SF \}$$

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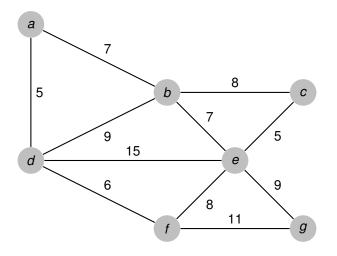
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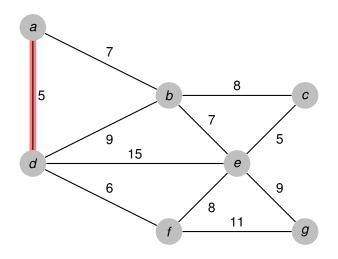
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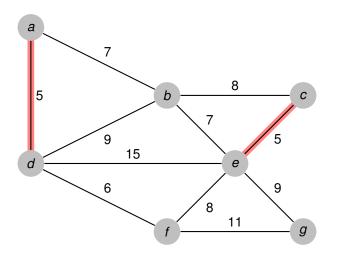
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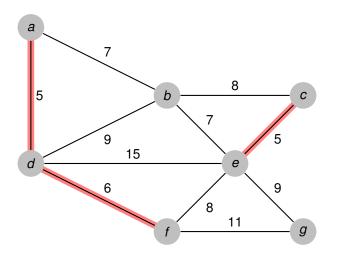
So $\{u, v\}$ is the edge with minimum weight among those crossing the cut.

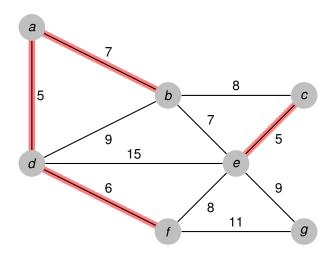


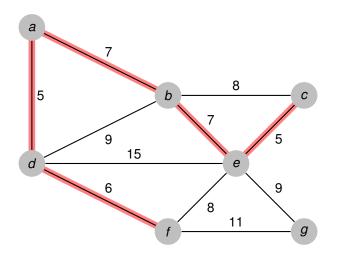
 $\textbf{Font:} \ \texttt{www.texample.net/tikz/examples/author/kjell-magne-fauske}$

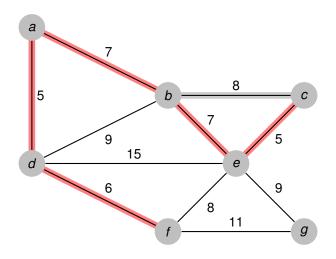


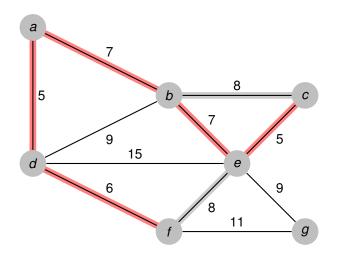


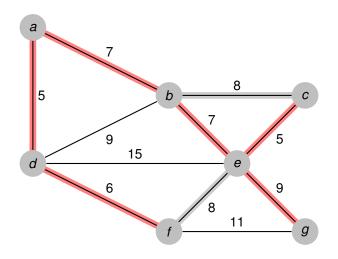












```
typedef pair < double, pair <int, int > WEdge;
typedef vector< vector< pair<double, int> > > WGraph;
int Kruskal(const WGraph& G, vector<pair<int,int>>& MST) {
  int n = G.size();
  UnionFind uf(n);
  int totalWeight = 0;
  priority_queue<WEdge, vector<WEdge>, greater<WEdge>> Q;
  for (int v = 0; v < n; ++v)
    for (auto& x: G[v]) if (v < x.second)
       Q.push({x.first, {v,x.second}});
  int sz = 0;
  while (sz < n - 1) {
    int u = Q.top().second.first;
    int v = Q.top().second.second;
    double wEdge = Q.top().first; Q.pop();
    if (uf.find(u) != uf.find(v)) {
      totalWeight += wEdge; ++sz;
      MST.push\_back(\{u,v\}); uf.merge(u,v);
  return totalWeight; }
```

```
class UnionFind {
   vector<int> rep;
   vector<int> size; // only valid for representative elements
   vector<vector<int>> elementsInClass; // only valid for reps
public:
   UnionFind(int n) : rep(n), size(n), elementsInClass(n) {
     for (int i = 0; i < n; ++i)
       \{rep[i] = i; size[i] = 1; elementsInClass[i] = \{i\}; \}
   int find (int e) {return rep[e];}
   void merge(int e1, int e2) {
     int r1 = find(e1); int r2 = find(e2);
     if (size[r1] > size[r2]) { // r1 new rep}
       size[r1] += size[r2];
       for (auto& x: elementsInClass[r2])
         {rep[x] = r1; elementsInClass[r1].push_back(x);} }
     else { // r2 new rep
       size[r2] += size[r1];
       for (auto& x: elementsInClass[r1])
         {rep[x] = r2; elementsInClass[r2].push_back(x); } }
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Let us consider the contribution to the cost of:

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Adding m edges to Q takes $O(m \log m)$ time.

Proof (continued):

Let us consider the contribution to the cost of the loop:

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int sz = 0;
while (sz < n - 1) {
  int u = Q.top().second.first;
  int v = Q.top().second.second;
  double wEdge = Q.top().first; Q.pop();
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 At each iteration we pop one element from Q,
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All in all, the cost is $O(m \log m)$ plus the cost of O(m) Finds and O(n) Merges.

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Find: costs constant time

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The total time is hence $O(m \log m) + O(m) + O(n \log n)$.

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Hence, Kruskal's Algorithm runs in $O(m \log n)$.