Design and Analysis of Algorithms

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- Two ideas are gleaming on the jeweler's velvet.
- First is the *calculus*, the second, the *algorithm*.
- The calculus made modern science possible; but it has been the algorithm that has made possible the modern world.
- David Berlinski, The Advent of the Algorithm.

The idea of the algorithm is at least 300 years old, so its understanding requires much work and effort.

The main goal of this class is to deepen our thinking about algorithms.

This is necessary to excel in your profession, but, perhaps more importantly, it can substantially improve your thinking skills. Basic computation - systematic manipulation of input data.

Description of such a manipulation is called *algorithm*.

An algorithm presupposes a fixed collection of elementary data types and operations which can be directly executed by a *computer* (a person or a machine).

Computation is normally used to obtain output data which satisfy some desired properties (often referred to as *specification*). *Given*: N men and N women. Each man ranks all the women (no equal rank is allowed). Similarly for each women.

Problem: Find an engagement between men and women such that

- Engagement is between one man and one women.
- Everyone must be engaged.
- Engagement is stable, i.e. there are no pairs $\langle m_1, w_1 \rangle$ and $\langle m_2, w_2 \rangle$ such that
 - m_1 prefers w_2 to w_1 and
 - w_2 prefers m_1 to m_2 .

It is often useful to replace original problem by its mathematical version, often referred to as *model* of the problem.

In our case the model is simple: we have two sets of points and the corresponding preference relations.

Need to find one-to-one correspondence between the sets which satisfy stable condition on preferences.

Recall, that function $F:A\to B$ is one-to-one correspondence if

- Different elements of A are mapped into different elements of B.
- For every element Y of B there is an element X in A such that Y = F(X).

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We use variable \mathfrak{m} for men and w for women.

function match

while $\exists m (free_man(m) \land \exists w \neg proposed(m, w)) do$

- **1** Select a free man m.
- It m propose to highest-ranked woman w he has not yet proposed.
- \bigcirc if w is free let her accept, otherwise
- if w prefers m to her current fiance let her break the engagement and accept m's proposal.

return the set of all engaged pairs.

Correctness Proof

1. The algorithm terminates. At every step a man proposes to a woman he has not proposed yet. Therefore, the maximum number of steps is $N \times N$.

2. The algorithm defines one-to-one correspondence between men and women.

- A man is engaged to at most one woman. A man proposes (and hence can get engaged) only if he is free.
- Every man is engaged. Suppose m is not engaged, i.e. he was rejected by every woman. A woman rejects a proposal only if she is engaged, and once engaged she stays engaged. Thus, at the end every woman is engaged to some man. Since a man cannot be engaged to two women (see above), N men are engaged. Contradicts our assumption.

• Different men are engaged to different women. A woman remains engaged from the point at which she receives her first proposal, and, since ties are not allowed, every change increases the quality of her partner. Hence she cannot be engaged to two different men.

This completes the proof of (2).

Correctness proof

3. The final engagement S is stable. Suppose it is not, i.e. S contains pairs (m_1, w_1) and (m_2, w_2) such that

- m_1 prefers w_2 to w_1 and
- w_2 prefers m_1 to m_2

To get engaged to w_1 , m_1 had to propose to her at some iteration. Since m_1 prefers w_2 he had to propose to w_2 at earlier iteration. There are two possible outcomes:

(a) m_1 and w_2 became engaged. To see that this is impossible let us notice that every change of engagement for a women increases quality of her partner so she would never get engaged to m_2 . Contradiction.

(b) w_2 was engaged to someone better than m_1 . By the same argument she would never get engaged to m_2 . Contradiction. m_1 could not have been engaged to $w_1!$ First considered by David Gale and Lloyd Shapley (two mathematical economists), in "College Admission and Stability of Marriage", 1962:

Given a set of preferences among employers and applicants, can we assign applicants to employers so that for every employer E, and every applicant A who is not scheduled to work for E, at least one of the following holds:

- **Q** E prefers everyone of its accepted applicants to A; or
- **2** A prefers his current situation over working for E.

In 2012 Shapley received a Nobel Prize in Economics: "for the theory of stable allocations and the practice of market design".

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First there could be more than one stable matching.

Example:

- \mathfrak{m} prefers w to w'
- \mathfrak{m}' prefers w' to w
- w prefers m' to m
- w' prefers m to m'

Our algorithm returns $\langle \mathfrak{m}, w \rangle$ and $\langle \mathfrak{m}', w' \rangle$

But, $\langle \mathfrak{m}, \mathfrak{w}' \rangle$ and $\langle \mathfrak{m}', \mathfrak{w} \rangle$ is also a stable matching!

It will not be returned by our algorithm no matter which free man is selected first. It does not! First, some definitions:

w is a *valid partner* of \mathfrak{m} if $\langle \mathfrak{m}, w \rangle$ belongs to some stable matching.

w is the *best valid partner* of m (best(m)) if w is a valid partner of m and no woman ranked higher than w is a valid partner of m.

Let $S^* = \{ \langle m, best(m) \rangle \}$

1. Suppose this is not the case, i.e. there is a run I of the algorithm which returns $S_I \neq S^*$.

2. Then there is a man who, in run I, proposed to his valid partner and was (immediately or eventually) rejected by her.

3. Let m be the *first* such man, and denote the woman he proposed to by w.

4. Since everyone is engaged, and a woman rejects a proposal only if she gets a better partner, S_I contains $\langle m', w \rangle$ where w prefers m' to m.

Theorem: Every execution of our algorithm returns S*

5. Since w is a valid partner of m, there is stable matching S such that

 $\langle \mathfrak{m}, \mathfrak{w} \rangle \in S.$

In S, m' is engaged to some woman. Let us denote her by w', i.e.

 $\langle \mathfrak{m}', \mathfrak{w}' \rangle \in S$

We have already shown that w prefers m' to m. But who does m' prefer, w or w'?

6. Since m is the *first* man rejected by his valid partner in run I, m' has not been rejected by any valid partner when he got engaged to w, i.e. he prefers w to w'.

7. Therefore, S is unstable. Contradiction!

Let α be an algorithm defined on a set D of strings, and T(n) be the worst-case running time of α on input of size n.

T(n) is order f(n) (T(n) is O(f(n))) if there are constants C and x_0 such that for every $x > x_0$, $T(x) \le C \times f(x)$.

We also say that f is an asymptotic upper bound of T. Asymptotic lower bound, g, of T is defined similarly and denoted by $\Omega(T)$.

Show that $T(n) = n^2 + 5n$ is in $O(n^2)$.

A deterministic algorithm is *polynomial* if it has an asymptotic polynomial upper bound.

There is one resource which can be used by at most one person at a time.

Multiple people request the use of this resource. A request is of the form "Can I use the resource starting at time s, until time f?"

The goal of a scheduler is to maximize the number of requests accepted.

Given: A finite set $R=\{i_1\ldots,i_n\}$ of names. Elements of R are called *requests*. Each request i is associated with an interval [s(i),f(i)).

A *schedule* of R is a subset of R containing no overlapping request.

Find: A best (optimal) schedule. i.e. schedule A of R such that no schedule of R contains more requests than A.

function schedule(R_0 : set of requests) : set of requests % returns a best schedule for R_0 . var R.A : set of requests

$$R := R_0, \qquad A := \emptyset.$$

while $R \neq \emptyset$ do

1. select a request $i \in R$ with earliest finish time,

2. add i to A,

3. remove from R all requests which overlap with i (including i). return A. In our scheduling algorithm we have:

- A *candidate set*, R from which a solution A is created.
- A selection function (earliest finish time) which, at each step, chooses the candidate to be added to the potential solution.

A greedy algorithm follows its selection function, never reverses its coices, and hopes for the best.

It is your job to guarantee that the hope is justified and a truly best solution is found.

Lemma 1. The algorithm terminates.

Termination follows since during each step R decreases in size.

In what follows, by A(m) and R(m) we denote the values of A and R after the execution of the m'th step of the loop.

Lemma 2. The algorithm returns a schedule.

Show by induction on k that for every step k:

(a) Intervals in A(k) are pairwise disjoint.

(b) No interval in R(k) overlaps with an interval in A(k). Base. After the first execution $A(1) = \{i_1\}$ and, due to step (3), no interval in R(1) overlaps with i_1 (step 3).

Inductive step: Suppose (a) and (b) hold for k = m.

 $A(m + 1) = A(m) \cup \{i_{m+1}\}$ where $\{i_{m+1}\}$ is selected from R_m . By inductive hypothesis intervals in A_m are pairwise disjoint and no interval in R(m) overlaps with an interval in A(m). Hence, intervals in A(m + 1) are pairwise disjoint, and, by step (3) of the algorithm, (b) holds for k = m + 1. Lemma 3. Let $O = \{j_1, \ldots, j_n\}$ be an optimal solution of the problem and A be returned by our algorithm. Then,

- $|A| \leq |O|;$
- For every step k of the algorithm

 $(*) \quad f(\mathfrak{i}_k) \leq f(\mathfrak{j}_k).$

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The first condition follows immediately since O is optimal and A is a solution (Lemma 2). Let us prove the second condition by induction on k. Base: k = 1 - is true by construction. Inductive Hypothesis: (*) holds for every $k \le m$. Show that (*) holds for k = m + 1.

Since O is a schedule we have that

 $(1) \quad f(j_m) \leq s(j_{m+1}).$

By inductive hypothesis,

 $(2) \ f(\mathfrak{i}_{\mathfrak{m}}) \leq f(\mathfrak{j}_{\mathfrak{m}}).$

(1) and (2) imply

 $(3) \ f(\mathfrak{i}_m) \leq s(\mathfrak{j}_{m+1})$

This means that at the moment of selection of i_{m+1} interval j_{m+1} does not overlap with intervals in A, and therefore belongs to R. Since the algorithm selects interval from R with the earliest finish time, we have

 $(4) \quad f(\mathfrak{i}_{m+1}) \leq f(\mathfrak{j}_{m+1})$

and (*) follows by induction.

Correctness

Lemma 4. A is optimal.

Suppose it is not and there is optimal $O = \{j_1, \ldots, j_n\}$, i.e. O is a schedule and |O| > |A|. Let i_m be the interval added to A during the last iteration. By Lemma 3,

 $(1) \quad f(i_m) \leq f(j_m).$

Since |O| > |A|, there is $j_{m+1} \in O$. Since O is a schedule,

 $(2) \quad f(j_m) \leq s(j_{m+1}).$

and, by (1) and (2)

 $(3) \quad f(\mathfrak{i}_m) \leq s(\mathfrak{j}_{m+1}).$

i.e. after the last iteration of the algorithm $j_{m+1}\in R.$ But, at this time, R must be empty. Contradiction.

Lemmas 2, 4 imply algorithm's correcteness.

Our scheduling algorithm traverses R twice (in steps (1) and (3)) and hence its time complexity is n^2 .

Can we refine the algorithm to make its time complexity be $O(n * \log n)$?

Step one can be reduced to sorting R and traversing it from left to right which is $O(n * \log n)$.

Instead of removing from R all intervals overlapping with the just scheduled interval i we can simply traverse R in search of intervals to be added to A.

This can be done as follows:

Let R[1..n], A[1..n] be arrays of requests where R[k] is a record containing k'th starting and finishing times.

1. Sort R in order of finishing time. [O($n * \log n$) time] 2. k,i := 0

- 3. f := -1 % finishing time of the last scheduled request
- 4. while k < n do [Linear Time]
- 5. k := k + 1
- $\textbf{6.} \qquad \textbf{if } R[k].s \geq f \textbf{ then }$

i := i + 1A[i] := R[k]f := R[k].f

7. return A[1..i].

GIVEN: A large set of identical resources and a set $R = \{i_1 \dots, i_n\}$ of requests.

Each request i is associated with an interval [s(i), f(i)).

A request i asks for the use of one resource during the interval [s(i), f(i)).

DO: Schedule *all* requests using as *few* resources as possible.

Examples: Schedule classes using as few classrooms as possible, allocate printing jobs using as few printers as possible, etc.

The set R of requests (also called jobs) is defined as in Problem 2. Resources are numbers from 1 to n.

To define a schedule we need the following notation: Let

$$\alpha: \mathbf{R} \to [1..n]$$

be an assgnment of jobs to resources and

$$R_k = \{j \in R: \alpha(j) = k\}$$

be the set of all jobs served by a resource k.

Mapping $\alpha : R \to [1..n]$ is called a *schedule* if

- \bullet For some $0 < d \leq n, \ R_1, \ldots, R_d$ is a partition of R, i.e.
 - $R=R_1\cup\cdots\cup R_d$ and
 - if $m \neq k$ then $R_m \cap R_k = \emptyset$.
- For every $0 < m \le d$, intervals requested by jobs from R_m do not overlap.

Schedule which uses as few resources as possible is called *optimal*.

depth(R) - max number of overlapping requests in R.

Can we partition R into R_1, \ldots, R_d where d = depth(R)and requests in any R_i do not overlap?

If the answer is positive then this allocation of resources and the subsequent scheduling of jobs is optimal – clearly, the number of resources needed is at least d. IDEA:

1. Schedule jobs according to their starting times.

2. Divide the set of available resources into those which are in use (*occupied*), had been used but are now available (*released*), and not being used at all. If possible, select next resource from *released*.

Pseudocode

function IntervalPartitioning(R : set of jobs) % Returns an optimal schedule (R_1, \ldots, R_m) for R. var released, occupied : resources; m : resource m := 0; released, occupied $:= \emptyset;$

- 1. Sort R w.r.t. starting times of its jobs.
- **2.** for every $I \in R$ do
 - 3. Move all resources in occupied which finished before the start of I into released.
 - 4. if released $\neq \emptyset$ then

m := select(released);

Move m from released to occupied.

- 5. else m := m + 1; % select new resource.
 - create R_m , set it to \emptyset , add m to occupied.
- 6. Add I to R_m .

Let jobs in R_i be served by resource i.

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- Algorithm obviously terminates after all jobs are scheduled.
- Algorithm returns a schedule.
- 1. Since every $I \in R$ is moved into some R_i (step 6)
- $\mathbf{R}=\mathbf{R}_1\cup\cdots\cup\mathbf{R}_{\mathfrak{m}}.$

No job is served by two resources. $R_1\cup\cdots\cup R_m$ is a partition.

2. No two jobs in $R_{\rm i}$ overlap. [Induction on number of iterations]

Note that when resource m is allocated to serve I every job served by resources in *released* finishes before I starts.

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To show that the schedule $R_1,\ldots R_m$ is optimal, it suffices to show that $m\leq d.$

Suppose (3) is not the case.

Then, before the allocation, every resource from 1 to d is in *occupied*, i.e. every such resource is in use. But this means that there are at least d + 1 overlapping requests in R which contradicts the definition of d.

Since any solution needs at least d resources, m = d, and R_1, \ldots, R_m created by the algorithm is an optimal solution.

GIVEN: Set R of requests, where each $i \in R$ has duration, t_i , and deadline, d_i . A single, but unlimited, resource, e.g. a processor.

Schedule maps a request $i\in R$ into interval [s(i),f(i)) of length t_i such that intervals for different requests are disjoint.

A request i is *late* if $f(i) > d_i$. A request's *lateness*, L_i is $f(i) - d_i$ if i is late and 0 otherwise.

DO: Find a schedule which

- starts at a given point s;
- minimizes maximum lateness, $L = max_i(L_i)$.

Basic Idea: schedule request with earliest deadline first!

min-max-lateness(R : set of requests, s : time) : schedule var f : time % finishing time of the last scheduled job.

Sort requests in order of their deadlines: $d_1 \leq \cdots \leq d_n$. f := s; For every job i from 1 to n do s(i) := f $f(i) := s(i) + t_i$ $f := f + t_i$ return [s(i), f(i)) for every i

return [s(i), f(i)) for every i.
idle time - time between jobs.

inversion - assignment of times to two jobs in which the job with later deadline is scheduled first, i.e.

- s(i) < s(j),
- $\bullet \ d_j < d_i$

Obviously, the schedule produced by our algorithm has no idle time and no inversion. Optimality follows from the following Lemmas:

Lemma 1. All schedules with no idle time and no inversion have the same maximum lateness.

Lemma 2. There is an optimal schedule with no idle time and no inversion.

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Let us compute the maximum lateness.

First notice, that two different schedules with neither inversion nor idle time only differ in order in which jobs with the same deadline are scheduled.

Clearly, all rearrangement of jobs with the same deadline start and finish at the same time, say s_d and f_d . Hence the maximum lateness for every rearrangement is the same $-f_d - d$.

Let O be an optimal schedule. Clearly it has no idle time.

Suppose it has an inversion. Then there are two consecutively scheduled jobs, i and j such that

(1) $d_j < d_i$.

Let us swap i and j and show that the maximum lateness, \overline{L} of the new schedule, \overline{O} does not exceed the maximum lateness, L of O, i.e. \overline{O} is still optimal.

The optimal schedule without inversion and idle time can be, therefore, obtained by repeating swapping until all inverse pairs are eliminated.

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Proof of Lemma 2

To show that $\overline{L} \leq L$ consider a job r in O scheduled for interval [s(r), f(r)) with lateness L_r .

The corresponding quantities in \bar{O} will be denoted by $[\bar{s}(r), \bar{f}(r))$, and \bar{L}_r .

The only possible increase in lateness \overline{L} in \overline{O} could have occur because of increase in lateness of i. Let us show that this is impossible. First, by definition, $\overline{f}(i) = f(j)$ and hence

$$(2) \ \bar{L}_i = \bar{f}(i) - d_i = f(j) - d_i.$$

From (1) and (2) we have

$$(3) \ \bar{L}_i = f(j) - d_i \ < \ f(j) - d_j = L_j.$$

Therefore,

$$\bar{L}_i < L_j \leq L$$

Spanning Trees

Given: Graph $G=\langle V,E\rangle$ with positive cost c_e associated with every edge e.

Definitions:

- An undirected graph is a *tree* if it is connected and has no cycles (recall that a cycle must include at least two different links).
- Spanning tree of G is a set T of edges such that $\langle V,T\rangle$ is a tree.
- Cost of T is the sum of costs of its edges.
- *Minimum spanning tree* of G is a spanning tree of G with minimal cost.

Problem: Find a minimum spanning tree of G.

Number of spanning trees in a graph is $O(n^n)$. Exhaustive search is impossible. kruscal($G = \langle V, E \rangle$: weighted graph) : set of edges % Assumption: all costs are different % returns a minimal spanning tree of G.

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var T : set of edges
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 $T := \emptyset$ Sort E in increasing order of costs. for every $e \in E$ do if $T \cup \{e\}$ has no circles then $T := T \cup \{e\}$ return(**T**). *Cut Property.* If S is a non-empty set of nodes different from V and $e = \langle v, w \rangle$ is a minimum-cost edge with one end in S and another in $V \setminus S$ then every minimum spanning tree contains *e*.

Proof. Let e be as above. Suppose there is a minimum spanning tree \top not containing e.

Since $\langle V, T \rangle$ is a tree, ν and w are connected by a path $P = \nu, \dots, \nu_1, w_1, \dots w$ in G where $e_1 = \langle \nu_1, w_1 \rangle$ is the first edge in P with one end in S and another in $V \setminus S$.

Let T_1 be obtained from T by replacing e_1 by e.

We will show that T_1 is a spanning tree of G. Since cost of T_1 is smaller than that of T this will contradict the minimality of T.

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To show that T_1 is a spanning tree we need to show that (a) Graph $\langle V, T_1 \rangle$ is connected.

Take two nodes x_1 and x_2 of V. Since T is a spanning tree, $\langle V, T \rangle$ is connected, i.e. contains a path Q connecting these nodes. If Q does not contain e_1 it is also a path of $\langle V, T_1 \rangle$ (which connects x_1 and x_2). If Q contains $e_1 = \langle v_1, w_1 \rangle$, i.e. $Q = \langle x_1, \dots, v_1, w_1, \dots, x_2 \rangle$ then path $R = \langle x_1, \dots, v_1, \dots, v, w, \dots, w_1, \dots, x_2 \rangle$ in T_1 connects x_1 and x_2 .

(b) $\langle V, T_1 \rangle$ has no cycles.

The only cycle in $\langle V, T_1 \cup \{e_1\}\rangle$ is the one composed of e and P. But this cycle can not be in $\langle V, T_1 \rangle$ since e_1 has been removed.

By the Cut Property every edge added to T in the Kruskal's algorithm belongs to every minimum spanning tree of G.

So, if T is a spanning tree then it is also minimum spanning tree.

Show that it is a spanning tree, i.e. $\langle V, T \rangle$ is a tree. Clearly, T has no cycles. Suppose it is not connected i.e.

there is set S different from \emptyset and V such that nodes from S and V \ S are not connected by edges from T. But there is an edge from S to V \ S in G. Cheapest such edge would have been added to T by the algorithm. Contradiction.

Implementing Kruscal's Algorithm

Naive implementation of the algorithm may require $|E| \times |V|$ steps. Can we do better?

Challenge: How to check that $\mathsf{T} \cup \{e\}$ has no cycles in time better than $|\mathsf{V}|?$

Idea: Represent T as a collection T_{i_1}, \ldots, T_{i_k} of disjoint sets. Design function find(u) which, for every $u \in S$ returns i_m such that $u \in T_{i_m}$.

Let $\{e = \langle v, w \rangle\}$, m = find(v) and n = find(w), and $|T_{i_m}| \ge |T_{i_n}|$. $T \cup \{e\}$ has no cycles iff $m \ne n$. In this case replace T_{i_m}, T_{i_n} by $T_{i_m} = unite(T_{i_m}, T_{i_n})$.

Good data structure allow $\log(|V|)$ find and constant unite.

One of the early applications of graph theory to computing. Even though the graphs in some form appeared in math starting with Euler, the first textbook on graph theory was published in 1935.

Similar algorithm was first published by a Czech mathematician Boruvka (1926) in a paper "Contribution to the solution of a problem of economical construction of electrical networks (Czech)".

In 1954 graph theory was sufficiently known so Kruscal could publish his work in a mathematical journal.

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Now the algorithm forms the basis for solutions of many optimization problems. GIVEN: directed graph $G = \langle V, E \rangle$ and the length $l_e \ge 0$ for each edge e; starting node s.

FIND shortest path from s to each other node.

Length, l(P) of path P is the sum of the lengths of its edges.

 $\mathbf{fringe}(\mathbf{S})$ - set of all $X \in V \setminus S$ connected to some node of S by an edge.

Simplification: (a) determine the *length* of the shortest path from s to each other node. (b) Assume that there is a path from s to any other node in G.

Our goal is to "serve", i.e., compute minimal distance from s, for each vertice of G. To do that

1. Maintain $S \subseteq V$ such that for each $u \in S$ the shortest distance, d(u) from s to u had been already computed, and is less or equal to d(v) for every $v \notin S$.

2. Consider set $E=\{\langle u,v\rangle: u\in S, v\not\in S\}$ and function α defined on E such that

$$\alpha(\langle \mathfrak{u}, \mathfrak{v} \rangle) = \mathfrak{d}(\mathfrak{u}) + \mathfrak{l}(\langle \mathfrak{u}, \mathfrak{v} \rangle),$$

select $\langle u_0, v_0 \rangle$ which minimizes α , add v_0 to S, and set $d(v_0)$ to $\alpha(\langle u_0, v_0 \rangle)$.

Clearly, new S still satisfies properties from (1).

Dijkstra(G : directed graph, l - length, s : node) % returns function d(u) whose value is the shortest % distance between s and u.

var S : set of explored nodes; function $d(\mathfrak{u})$

 $S := \{s\} \qquad d(s) := 0$ while $S \neq V$ do 1. Select $v_0 \in Fringe(S)$ for which $d'(v) = \min_{\{u \ : \ u \in S, \langle u, v \rangle \in G\}} d(u) + l(\langle u, v \rangle)$ is as small as possible. 2. $d(v_0) := d'(v_0)$ 3. $S := S \cup \{v_0\}$ Potume d

Return d

How to represent G, S, and Fringe(S)?

- G adjacency list $O(max_degree(G))$.
- S a tree of nodes

An array, nodes, where nodes(v) contains a pointer to a position of v in S (or nil if v is not there yet).

F = Fringe(F) - a priority queue with entries $\langle v, key(v) \rangle$ where at each point of the execution of the algorithm key(v) is the length of a shortest path from s to v found so far. $S := \{s\}$ d(s) := 0

 $\mathsf{F} := \{ \langle \mathsf{y}, \mathsf{key}(\mathsf{y}) \rangle \}$

where y is adjacent to s and key(y) is the length of shortest edge from s to y.

while $F \neq \emptyset$ do 1. $m := extract_minimum(F)$ 2. $S := S \cup \{m\}$ 3. d(m) := key(m)4. For every edge $\langle m, v \rangle$ with $v \in V \setminus S$ do 4a. Temp $:= d(m) + l_{\langle m, v \rangle}$ 4b. if $v \notin F$ then insert(F, v, Temp) 4c. else if Temp < key(v) then change key(F, v, Temp)

return d

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A type of algorithm which breaks the input into parts, solves each part recursively, and then combines the solutions of these subproblems into an overal solution.

Example: computing the value of an arithmetic expression represented by a tree, merge sort, etc.

Problem: Given a person's preferences (for books, movies, etc.) match them with preferences of other people on the Web with similar interests to provide a suggestion.

Preferences are often defined by rankings, i.e., labeling the objects from 1 to n. So the problem is to *define and compute the distance between the rankings*. Given: a sequence $S = \langle a_1, ..., a_n \rangle$ of distinct numbers. Def: a pair i < j of indices form an *inversion* if $a_i > a_j$. Find: The number of inversions in S. Sequence $\langle 1, 2, 3 \rangle$ has no inversions, sequence $\langle 2, 1, 3 \rangle$ has

one, formed by indices 1 and 2, sequence (3,2,1) has three.

Basic Idea: Sort sequences together with counting inversions to facilitate merging.

 $\begin{array}{l} \mbox{CountInversions} \\ \mbox{Given: Sequence } S = \langle a_1, \ldots, a_n \rangle \\ \mbox{Return: Number of inversions in S and sorted S.} \\ \mbox{if } n = 1 \mbox{ then return}(0,S) \\ \mbox{Divide S into two halves, A and B.} \\ (r_A, A) := CountInversion(A) \\ (r_B, B) := CountInversion(B) \\ (r,S) := merge_and_count(A,B) \\ \mbox{return}(r_A + r_B + r,S). \end{array}$

Counting Inversions

Merge_and_count

Given: Sorted sequences A,B.

Return: The number of pairs X, Y such that $X \in A, Y \in B$ and X > Y, and sorted sequence C consisting of elements of A and B.

var i,j : ponter, C : sequence, Count : integer % i and j point to element a_i of A and element b_j of B i,j := 1 $C := \emptyset$ Count := 0 while both lists are non-empty do

Move the smaller of two elements a_i , b_i to C.

if $b_i < a_i$ then Count := Count + |A|.

Advance pointer to the list from which the smaller element was selected.

Append the remainder of the non-empty list to C. return (Count,C). Clearly, Merge_and_count requires O(n) where n=|A|+|B|.

Let T(n) denote the worst-case running time on input of size n. Let $n = 2^k$. To get the input down to 2 we need $k = \log_2 n$ levels of recursion.

First level has one merge which requires $c \times n$ steps.

On the i'th level the number of subproblems has doubled i times, so their number is 2^i . Each has input size $n/2^i$ and takes at most $c \times n / 2^i$ steps. Overall time spent on level i is $2^i \times (c \times n / 2^i) = c \times n$.

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Executing k levels requires $c \times n \times \log_2 n$, i.e. the algorithm time is $O(n \times \log_2 n)$.

- 1. Divide the problem into subproblems such that
 - The number of subproblems is polynomial.
 - The solution to the original problem can be easily computed from solutions to the subproblems.
 - Subproblems can be ordered from "smallest" to "largest" and solutions of larger are connected to that of smaller by a recurrence relation.

2. Get rid of recursion by storying solutions of problems from smaller to larger in an array.

Given: n requests, each request i associated with an interval $[s_i, f_i)$ and has the weight, v_i . One server.

Find: A subset of mutually compatible requests with maximum summary weight.

This time ordering intervals by the earliest finishing time does not work. Later interval not compatible with the selected one may have much higher weight.

First solve a simpler problem – find maximum summary weight.

Notation: $f_1 \leq f_2 \cdots \leq f_n$; p(j) is the last interval in the sequence which finishes before j starts, 0 if no such interval exists.

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Recursive Algorithm

Divide the problem into finding maximum weight of a schedule containing j and that not containing j.

opt(j:requests):weight

% Requests from 1 to j are sorted by finishing time.

% Returns weight of the optimal solution of the

- % scheduling problem.
- % Recurrence relation:

$$opt(j) = max(v_j + opt(p(j)), opt(j-1))$$

if j = 0 return(0). return(max(v_j + opt(p(j)), opt(j - 1))).

% Returns max of the best solution containing j % and that not containing j.

Unfortunately has exponential number of calls. Can we reduce this number?

The first time opt(k) is computed store the value in an array, say M[0...n].

```
opt(n:requests): array
% Requests 1...n sorted by finish time
% Returns M[0..n] where M[j] is the weight of
% optimal scheduling of first j requests.
var M[0..n].
M[0] := 0
for j from 1 to n do
```

```
M[j] := \max(\nu_j + M[p(j)], M[j-1])
return(M)
```

Find_Solution(M : array, j : index) : sequence of requests.

% For j>0, M[j] contains the weight of optimal solution % for scheduling first j requests.

% Returns optimal solution for scheduling first j % requests.

The function is in O(n).

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Given: A set $S = \{(x_1, y_1) \dots (x_n, y_n)\}$ of points on the coordinate plane Find a segment of best fit.

How to define *best fit*, i.e., how to measure an error – distance between S and function f(x, a, b) = ax + b is an interesting theoretical question.

A popular *least squares* method finds a and b which minimize

$$R^2 = \sum_{i=1}^n (y_i - f(x_i, a, b))^2$$

There are good algorithms for such a minimization, and we assume that they are given.

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What to do if S can only be reasonably approximated by more than one segment?

1. Partition S into a number of segments and approximate each segment separately.

2. Balance the number of segments in the approximation and the quality of the fit.

In other words, fit points well using as few segments as possible.

We order partitions by assigning *penalties* – the sum of the number of segments in the partition (times some constant C) AND the error value of the optimal line through each segment.

The first value keeps the number of segments small. The second ensures that each component of the partition is accurately approximated by a segment.

Segmented Least Square Problem: Given C, find a partition of minimum penalty.

Sort points p_1, \ldots, p_n with respect to X-coordinate. opt(n) penalty of optimal approximation of p_1, \ldots, p_n (0 if i = 0).

 e_{ij} - minimum error of any line approximating p_i, \ldots, p_j . If p_i, \ldots, p_n is the best last segment then

$$opt(n) = C + e_{in} + opt(i)$$

The recurrence relation

$$opt(j) = min_{1 \le i < j}(C + e_{ij} + opt(i))$$

opt(p₁,..., p_n: points): array
% Points are sorted by X coordinate.
% Returns M[0..n] where M[j] is the minimum penalty
% for approximation of p₁,..., p_j.
var M[0..n], e[0..n, 0..n]
M[0] := 0

for every $i \le j \le n$ do

compute minimal penalty e_{ij} for approximation of p_i, \ldots, p_j by one line. for j = 1 to n do

$$M[j] := \min_{1 \le i < j} (C + e_{ij} + M[i])$$

return(M)

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Given: Single machine to process n job requests. Machine is available in time interval from 0 to W. A request i requires time w_i to process.

Goal: find a schedule which keeps the machine as busy as possible.

Equivalent formulation: Given a set of items with non-negative weights w_i and integer W find a subset S of items such that

$$\sum_{i\in S} w_i \leq W$$

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• subject to restriction above the sum is as large as possible.

$\operatorname{opt}(\mathfrak{i},w)$ - solution of the problem with items $1\ldots\mathfrak{i}$ and allowed weight w.

Recurrence relation:

$$opt(i, w) = max(opt(i-1, w), w_i + opt(i-1, w-w_i))$$

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Subset Sum

function MaxWeight input: items $1 \dots n$, weights w_1, \dots, w_n , max weight W output: $M[i, w] = \max(\sum_{k \in S} w_k \le w \text{ where } S \subseteq \{1, \dots, i\})$ var : array M[0...n, 0...W]M := 0For every i do For every w do if $w < w_i$ then (ith item does not fit) M[i, w] := M[i - 1, w]else $M[i, w] := \max(M[i-1, w], w_i + M[i-1, w-w_i])$

return(M)

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To complete the solution of the original problem compute an array M, M := MaxWeight(P) and call function print(n,W) defined as follows:

```
function print(i, w)

if i > 0 then

if M[i, w] \neq M[i - 1, w] then

display(i)

print(i - 1, w - w<sub>i</sub>)

else print(i - 1, w)
```

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knapsack(n : item, W : capacity) : set of items % items 1..., each item i has weight w_i and value v_i % Returns subset of items which maximizes the sum of % their values subject to the capacity restriction.

function MaxVal(n : items, W : knapsack capacity) : int Recursive algorithm to compute maximim value of items which can be packed in a knapsack of capacity W.

$$if n = 0 return 0 \\ if W < w_n then return(MaxVal(n - 1, W)) \\ return(max(MaxVal(n - 1, W), v_n + MaxVal(n - 1, W - w_n)))$$

Knapsack (memoization)

MaxValues(n : items, W : knapsack capacity) : array %Create array M such that M[i, w] = MaxVal(i, w).

Set the entries of M to 0.

For every
$$i = 1$$
 to n do
For every $w = 1$ to W do
if $w < w_i$ then (ith item does not fit)

$$M[i,w] = M[i-1,w]$$

else

$$M[i, w] = max(M[i-1, w], v_i + M[i-1, w-w_i])$$

return(M)

To complete the solution of the original problem compute an array M, M := MaxValues(n, W) and call function print(n, W) defined as follows:

```
function print(i, w)

if i > 0 then

if w < w<sub>i</sub> then print(i - 1, w)

else

if M[i - 1, w] \ge v_i + M[i - 1, w - w_i] then

print(i - 1, w)

else display(i)

print(i - 1, w - w_i)
```

C(n,k) - number of subsets (combinations) of k-elements from an n-element set.

Standard formulas:

$$C(n,k) = n!/(k!(n-k)!)$$

$$C(n,k) = n \times n - 1 \times \cdots \times (n-k+1)/k!$$

and

$$C(n,k) = C(n-1,k-1) + C(n-1,k)$$
 for $0 < k < n$
$$C(n,0) = C(n,n) = 1$$

function comb(n, k)input: n > k > 0output: C(n, k)var matrix M[n, k] of integers for i from 0 to n do for j from 0 to min(i, k) do if j = 0 or j = i then M[i, j] := 1else M[i, j] := M[i - 1, j - 1] + M[i - 1, j]return(M)

A problem belongs to class NP if it can be solved in polynomial time by a non-deterministic Turing machine.

To solve a problem such a machine non-deterministically selects the problem's candidate solution and checks, if it is indeed a solution, in polynomial time.

Every problem belonging to a class P of polynomially solvable problems also belongs to class NP. It is not known if the reverse is true.

Figuring out if P = NP is one of the most important problems of CS.

Decision problem: Given a set L of string over alphabet Σ (often called a language) check if a string $x \in \Sigma^*$ belongs to L. Let $L_1, L_2 \subseteq \Sigma^*$.

A polynomial function $\tau: \Sigma^* \to \Sigma^*$ is a *polynomial reduction* of L_1 to L_2 if for every $x \in \Sigma^*$, $x \in L_1$ iff $\tau(x) \in L_2$.

A decision problem for L is NP-complete if

- L belongs to NP.
- Every NP problem is polynomially reducible to L.

It is easy to see that P = NP iff there is a polynomial solution of at least one NP-complete problem.

A *literal* is a boolean variable or its negation.

Clause is a set of literals. A clause C is *satisfied* by an assignment of truth values to variables if this assignment makes at least one elements of C true.

Formula is a set of clauses. Formula F is satisfied by a truth assignment I if all of its clauses are satisfied by I. Assignment I satisfying F is called a *model* of F.

Problem: Check if a given formula is satisfiable, i.e. has a model. (Here our language L consists of all satisfiable formulas).

SAT

It is easy to prove that SAT is an NP problem.

The corresponding non-deterministic Turing Machine will non-deterministically select an assignment

 $I = \{X_1 = \nu_1, \dots, X_n = \nu_n\} \text{ of the truth values to the variables of } F \text{ and checks if } I \text{ is a model of } F.$

The latter can be done in polynomial time as follows:

repeat

Select $X_i = v_i$.

Remove all clauses containing a literal ι formed by X_{ι}

which is true under this assignment.

Remove from the clauses of F all remaining

occurrences of such literals.

until $F = \emptyset \lor \{ \} \in F$.

if $F = \emptyset$ return true else return false.

In the early 70s of the last century Cook and Levin independently established that SAT is an NP- complete problem.

Despite the apparent absence of a polynomial algorithm for solving SAT its instances are frequently solved in many practical applications (including planning, diagnostics, decision support systems, etc.) by programs called SAT-solvers.

In the last 20 years SAT-solvers moved from solving problems with 100 variables and 200 clauses to $1,000,000^+$ variables and $5,000,000^+$ clauses.

Partial interpretation of a formula F is a mapping I of a subset of its variables to the truth values.

 $\begin{array}{l} \mbox{function SAT} \\ \mbox{input: Formula } F_0 \mbox{ and partial interpretation } I_0. \\ \mbox{output: } \langle I, true \rangle \mbox{ where } I \mbox{ is a model of } F_0 \mbox{ which extends } I_0. \\ & \quad \langle I_0, false \rangle \mbox{ if no such model exists.} \end{array}$

Uses function Cons(F, I) which expands I by the assignment of variables which must be true to satisfy F and simplifies F accordingly. If there are no expansions of I returns false.

var F: formula; I : partial interpretation; X : boolean.

$$\begin{split} F &:= F_0; \qquad I := I_0; \\ \text{if } Cons(F, I) = \text{false then} \\ & \text{return } \langle I_0, \text{false} \rangle. \\ \langle F, I \rangle &:= Cons(F, I). \\ \text{if } F &= \emptyset \text{ return } \langle I, \text{true} \rangle. \\ \text{select a boolean variable } p \text{ from } F \text{ undefined in } I. \\ \langle I, X \rangle &:= SAT(F \cup \{\{p\}\}, I). \\ \text{if } X = \text{true return } \langle I, X \rangle. \\ \text{return } SAT(F \cup \{\{\neg p\}\}, I). \end{split}$$

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function Cons(F, I)

Returns $\langle F', I'\rangle$ such that a model of F contains I iff it is a model of F' which contains I'.

Return *false* if no such F', I' exist.

while F contains a clause of the form {l} do Remove from F all clauses containing l. Remove from F all occurrences of \overline{l} where $\overline{p} = \neg p$, $\overline{\neg p} = p$. $I := I \cup \{l\}$ if $\emptyset \in F$ return false return $\langle F, I \rangle$. Let $I = \emptyset$ and $F = \{\{X_1\}, \{\neg X_1, X_2, X_3\}, \{\neg X_1, X_4\}\}.$

 $Cons(F,I) \text{ returns } F = \{\{X_2, X_3\}\} \text{ and } I = \{X_1, X_4\}$

Suppose the algorithm non-deterministically selects X_2 and calls SAT({{ X_2, X_3 }, { X_2 }}, { X_1, X_4 }).

The new call to SAT returns $I = \{X_1, X_4, X_2\}$ and $F = \emptyset$.

The second termination condition is satisfied and SAT returns $\langle \{X_1, X_4, X_2\}, true \rangle$.

Developers of SAT-solvers found many ways to improve solvers' efficiency, including

- Smart heuristics which allow selection of a good p.
- Learning clauses containing information about previous failures and adding it to the formula. This leads to a smart backtracking.
- Occasional random restarting of search to avoid being stuck in a wrong path.
- Smart data structurs.
- Automatic tuning of heuristics for particular instances.

SAT restricted to formulas whose clauses contain at most three literal is called 3-SAT.

To show that it is NP-complete we'll define a polynomial reduction τ from SAT to 3-SAT.

For every clause $C = \{\lambda_1, \dots, \lambda_n\}$, $\tau(C)$ consists of clauses

```
 \{ \lambda_{1}, \lambda_{2}, Y_{1} \}, \\ \{ \neg Y_{1}, \lambda_{3}, Y_{2} \}, \\ \dots \\ \{ \neg Y_{i-2}, \lambda_{i}, Y_{i-1} \}, \\ \dots \\ \{ \neg Y_{n-3}, \lambda_{n-1}, \lambda_{n} \} \}
```

where Ys are new variables.

Intuitively, Y_1 says that at least one λ_i where $i\geq 3$ must be true.

 $\{\neg Y_{i-2}, \lambda_i, Y_{i-1}\}$ says that if Y_{i-2} is true (i.e. at least one λ_k with $k \geq i$ is true) then λ_i or some literal with index greater than i is true.

Finally,

$$\tau(F) =_{def} \{\tau(C) : C \in F\}$$

Clearly, τ is polynomial.

To complete the proof we need to show that

 $F \in SAT \text{ iff } \tau(F) \in 3\text{-}SAT$

We divide the proof into two parts: (A) If $F \in SAT$ then $\tau(F) \in 3$ -SAT. (B) If $\tau(F) \in 3$ -SAT then $F \in SAT$. Suppose I is a model of F. Let C be an arbitrary clause of F. To define a satisfying assignment of $\tau(C)$ let us assume that λ_i is the first literal in C made true by I and expand I as follows:

- Set variables Y_1, \ldots, Y_{i-2} to true.
- Set variables Y_{i-1}, \ldots, Y_{n-3} to false.

Clearly, the result is a model of $\tau(C)$. Repeat the process for all C's from F. Since all the new variables in different clauses are different the resulting assignment is a model of $\tau(C)$.

Suppose I is a model of $\tau(F).$ It is not difficult to show that I is also a model of F.

Consider arbitrary $C\in F$ and show that at least one $\lambda_i\in C$ is satisfied by I.

Suppose it is not the case.

Then, to satisfy the first n-3 clauses of $\tau(C)$, I must set Y_1, \ldots, Y_{n-3} to true. But this would mean that the last clause is not satisfied. Contradiction.

The assumption is false, I satisfies C and, hence, F.

MaxSat: Given a set F of clauses and a natural number K check if there is a truth assignment which satisfies at least K clauses of F.

To show that MaxSat is NP-complete consider a mapping τ which maps a boolean formula F to a pair $\langle F, k \rangle$ where k is the number of clauses in F.

Clearly, τ is polynomial and $F \in SAT$ iff $\tau(F) \in MaxSat$.

 τ is a polynomial reduction and hence MaxSat is NP-complete.

Hamiltonian Cycle: Given a graph G is there a cycle which passes through each node of G exactly once?

Independent Set: Given an undirected graph G = (V, E) and an integer $K \ge 2$, is there a subset C of V such that $|C| \ge K$ and no two edges of C are connected by an edge from E?

3-coloring: Given an undirected graph G = (V, E) can we color the nodes in three different colors so that no nodes connected by an edge are of the same color? Among n identically looking coins one is fake, i.e. is lighter than a regular coin. Design an efficient algorithm to determine which coin is fake using a balance scale.

```
function fake(set S of coins)
% S contains one fake coin % return the fake coin
Divide S into two halfs, S_1 and S_2 with possibly one coin, c left
on the table.
If weight(S_1) = weight(S_2) then return(c).
If weight(S_1) < weight(S_2) then return(fake(weight(S_1)).
```

 $return(fake(weight(S_2))).$

W(n) – number of weighs needed to find the fake coin in a set of n coins.

$$W(1) = 0$$
$$W(n) = W(\lfloor n/2 \rfloor) + 1 \text{ for } n > 1$$

Solve for $n = 2^k$. $W(n) = W(2^k) = W(2^{k-1}) + 1 = W(2^{k-2}) + 1 + 1 = k + 1 = \log_2(n) + 1$ Check: $W(2 \times 2^{n-1}) = \log_2(2) + \log_2(2^{n-1}) = W(\lfloor n/2 \rfloor) + 1$ What if we divide into three parts, S_1, S_2, S_3 ?

1. No coin remains. Find the lightest, S (one weighing) and call $\mathsf{fake}(\mathsf{S})$

2. One coin, c, remains. If weight(S_1) - weight(S_2)

```
If weight(S_1) = weight(S_2) then
```

select c_1 from S_1

if weight(c_1) > weight(c) then return(c)

 $return(fake(S_3))$

etc.

Reccurrence Relation:

$$W(n) = W(\lfloor n/3 \rfloor) + 3$$
 for $n > 3$

Since $log_3 < log_2$ this method is faster. (8 versus 13 for n=10000)

Given: $p(X) = a_n X^n + a_{n-1} X^{n-1} + \dots a_0$ and the value, x of X. Efficiently compute: p(x).

A good representation of p(X) is obtained by successively taking X as a common factor from remaining polynomials of diminishing degrees (Horner Rule):

$$p(X) = 2X^{4} - X^{3} + 3X^{2} + X - 5 =$$

$$X(2X^{3} - X^{2} + 3X + 1) - 5 =$$

$$X(X(2X^{2} - X + 3) + 1) - 5 =$$

$$X(X(X(2X - 1) + 3) + 1) - 5 =$$

Substantially decreases the number of operations.

horner(P[0..n], x) P[0..n] = $(a_0, ..., a_n)$ V := P[0] for i from n - 1 to 0 do V := x × V + P[i]

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Let $n = b_k \dots b_0$ be a binary string representing a positive integer n and $p(x) = b_k x^k + \dots + b_i x^i + \dots + b_0$. Clearly, n = p(2)

Horner rule for p(2): V := 1for i from k - 1 to 0 do $V := 2V + b_i$ Computing $a^{p(2)}$: $a^V := a^1$ for i from k - 1 to 0 do $a^V := a^{2V+b_i}$

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$$\begin{split} a^{2V+b_i} &= a^{2V} \times a^{b_i} = (a^V)^2 \times a^{b_i} = \\ (a^V)^2 \text{ if } b_i &= 0; \ (a^V)^2 \times a \text{ otherwise} \\ V &:= a \\ \text{for i from } k-1 \text{ to } 0 \text{ do} \\ V &:= V \times V \\ \text{if } b_i &= 1 \text{ then } V := V \times a \\ \text{return}(V) \end{split}$$

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We use variable h for hospitals and s for student.

function match

while $\exists h (open_slot(h) \neq 0) do$

- Select a hospital h with an open slot.
- It h offer position to highest-ranked student s left on the list.
- if s is free let s accept and decrease open_slot(h)
 otherwise
- if s prefers h to h' to which s is committed let her break the engagement and accept h's proposal; decrease open_slot(h) and increase open_slot(h').

return the set of all engaged pairs.

Given: Spanning tree T with positive and distinct costs of edges.

Bottleneck of T is the edge with the greatest cost.

Question 1: Is every minimum-bottleneck spanning tree of G also a minimum spanning tree of G?

NO. Consider graph with nodes a, b, c, and d. Every pair of nodes is connected by an edge. c(a,b) = 8, c(a,c) = 6, c(b,d) = 10, c(c,d) = 12, c(c,b) = 2, c(a,d) = 11. Min-bottlneck = (c, a, b, d), Min-spanning = (d, c, b, a). Given: n minutes, two computers a and b, a_i - value of running machine a during minute i. Similarly for b_i .

Find: Use the computers to run your job to achieve maximum value. Constraints: Job can run on one machine at any given minute. A minute is required to change the machine.

opt(i:minutes) - best value for job allocation during first i minutes.

opt(i:minutes, C:computer) - best value for job allocation during first i minutes which ends in job being allocated to C.

$$opt(i) = max(opt(i, a), opt(i, b))$$

 $opt(i, a) = a_i + max(opt(i - 1, a), opt(i - 2, b))$

```
opt(n : min) : time

var M[0..n, a..b] : time

M[0, a], M[0, b] := 0

M[1, a] := a_1

M[1, b] := b_1

For i = 2 to n do

M[i, a] := a_i + max(M[i - 1, a], M[i - 2, b])

M[i, b] := b_i + max(M[i - 1, b], M[i - 2, a])

return max(M[n, a], M[n, b])
```

```
\begin{split} & \text{output\_plan}(M) \\ & \% \text{ outputs optimal allocation for $n$ minutes, $n \geq 0$} \\ & \text{if $M[n, a] > M[n, b]$ then $out(M,n,a)$} \\ & \text{else out}(M,n,b) \end{split}
```

```
out(M, i, C)
```

% outputs optimal allocation for i minutes which ends in % machine C; \bar{C} stands for "the other machine".

if i > 0 then print(C, at, i)

```
if i > 1 then
```

```
if M[i-1,C]>M[i-2,\bar{C}] then out(M,i-1,C) else out(M,i-2,\bar{C})
```

Given: a sequence $\langle s_1, \ldots, s_n \rangle$ where i is a week and s_i is a number of pounds to be shiped in this week.

Do: For every i schedule company A or company B to ship s_i . $cost(A, i) = r \times s_i$ cost(B, i) = c

Constraints: A contract with B must be made in blocks of four consequitive weeks. Ensure minimal cost.
To find minimal cost divide the problem of scheduling shipments for i weeks into two. In one, the last shipment is done by A, in the other one – by B.

Recurrence relation:

if i - 4 > 0 then

 $\texttt{opt}(i) = \texttt{min}(\texttt{opt}(i-1) + r \times s_i, \texttt{opt}(i-4) + 4 \times c)$

else

$$opt(i) = r \times s_1 + \cdots + r \times s_i$$
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function opt(i : index) : cost

var M[0..i].

M[0] := 0

for j from 1 to i do

if j - 4 > 0 then

M[j] := min(M[j - 1] + r \times s_j, M(j - 4) + 4 \times c)

etc

return(M)
```