

Algorithms: COMP3121/3821/9101/9801

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7. DYNAMIC PROGRAMMING



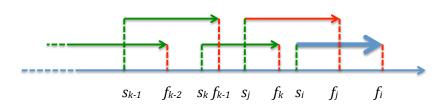
Dynamic Programming

- The main idea of Dynamic Programming: build an optimal solution to the problem from optimal solutions for (carefully chosen) subproblems;
- Subproblems are chosen in a way which allows recursive construction of optimal solutions to problems from optimal solutions to smaller size problems.
- Efficiency of DP comes from the fact that that the sets of subproblems needed to solve larger problems heavily overlap; each subproblem is solved only once and its solution is stored in a table for multiple use for solving many larger problems.

- Instance: A list of activities a_i , $1 \le i \le n$ with starting times s_i and finishing times f_i . No two activities can take place simultaneously.
- Task: Find a subset of compatible activities of maximal total duration.
- Remember, we used the Greedy Method to solve a somewhat similar problem
 of finding a subset with the largest possible number of compatible
 activities, but the Greedy Method does not work for the new problem.
- We start by sorting these activities by their finishing time into a non-decreasing sequence, so will assume that $f_1 \leq f_2 \leq \ldots \leq f_n$.
- For every $i \leq n$ we solve the following subproblems: **Subproblem** P(i): find a subsequence σ_i of the sequence of activities $S_i = \langle a_1, a_2, \dots, a_i \rangle$ such that:
 - \bullet σ_i consists of non-overlapping activities;
 - \circ σ_i ends with activity a_i ;
 - 3 σ_i is of maximal total duration among all subsequences of S_i which satisfy 1 and 2.
- Note: the role of condition 2 is to simplify recursion.

- Let T(i) be the total duration of the optimal solution S(i) of the subproblem P(i).
- For S(1) we choose a_1 ; thus $T(1) = f_1 s_1$;
- Recursion: assuming that we have solved subproblems for all j < i and stored them in a table, we let

$$T(i) = \max\{T(j) + f_i - s_i : j < i \& f_j < s_i\}$$



ullet In the table, besides T(i), we also store j for which the above max is achieved.



- Why does such a recursion produce optimal solutions to subproblems P(i)?
- Let the optimal solution of subproblem P(i) be the sequence $S = (a_{k_1}, a_{k_2}, \dots, a_{k_{m-1}}, a_{k_m})$ where $k_m = i$;
- We claim: the truncated subsequence $S' = (a_{k_1}, a_{k_2}, \dots, a_{k_{m-1}})$ is an optimal solution to subproblem $P(k_{m-1})$, where $k_{m-1} < i$.
- Why? We apply the same "cut and paste" argument which we used to prove the optimality of the greedy solutions!
- If there were a sequence S^* of a larger total duration than the duration of sequence S' and also ending with activity $a_{k_{m-1}}$, we could obtain a sequence \hat{S} by extending the sequence S^* with activity a_{k_m} and obtain a solution for subproblem P(i) with a longer total duration than the total duration of sequence S, contradicting the optimality of S.

• Continuing with the solution of the problem, we now let

$$T_{max} = \max\{T(i) : i \le n\}.$$

- We can now reconstruct the optimal sequence which solves our problem from the table of partial solutions, because in the i^{th} slot of the table, besides T(i), we also store j such that the optimal solution of P(i) extends the optimal solution of subproblem P(j).
- Why is such solution optimal?
- If such optimal solution ends with activity a_k , it would have been obtained as the optimal solution of problem P(k).
- Time complexity: having sorted the activities by their finishing times in time $O(n \log n)$, we need to examine n intervals in the role of the last activity in an optimal sub-sequence and for each such interval we have to find all preceding compatible intervals and their optimal solutions (to be looked up in a table). Thus, the time complexity is $O(n^2)$.

- Longest Increasing Subsequence: Given a sequence of n real numbers A[1..n], determine a subsequence (not necessarily contiguous) of maximum length in which the values in the subsequence are strictly increasing.
- Solution: For each $i \leq n$ we solve the following subproblems:
- Subproblem P(i): Find a subsequence of the sequence A[1..i] of maximum length in which the values are strictly increasing and which ends with A[i].
- Recursion: Assume we have solved all the subproblems for j < i;
- We now look for all A[m] such that m < i and such that A[m] < A[i];
- Among those we pick m which produced the longest increasing subsequence ending with A[m] and extend it with A[i] to obtain the longest increasing subsequence which ends with A[i];
- Why does this produce optimal solutions to subproblems? The same "cut and paste argument:

- Truncating optimal solution for P(i) must produce optimal solution of the subproblem P(m), otherwise we could find a better solution for P(i) as well, by extending such better solution of P(m).
- We store in the i^{th} slot of the table the length of the longest increasing subsequence ending with A[i] and m such that the optimal solution for P(i) extends the optimal solution for P(m).
- So, we have found for every $i \leq n$ the longest increasing subsequence of the sequence A[1..i] which ends with A[i].
- Finally, from all such subsequences we pick the longest one.
- Again, the condition for the sequence to end with A[i] is not restrictive because if the optimal solution ends with some A[m], it would have been constructed as the solution for P(m).
- Time complexity: $O(n^2)$.
- Exercise: (somewhat tough, but very useful) Design an algorithm for solving this problem which runs in time $n \log n$.

- Making Change. You are given n types of coin denominations of values v(1) < v(2) < ... < v(n) (all integers). Assume v(1) = 1, so that you can always make change for any integer amount. Give an algorithm which makes change for any given integer amount C with as few coins as possible, assuming that you have an unlimited supply of coins of each denomination.
- Solution: DP Recursion on the amount C. We will fill a table containing C many slots, so that an optimal solution for an amount i is stored in slot i.
- If C=1 the solution is trivial: just use one coin of denomination v(1)=1;
- Assume we have found optimal solutions for every amount j < i and now want to find an optimal solution for amount i.

- We consider optimal solutions opt(i-v(k)) for every amount of the form i-v(k), where k ranges from 1 to n. (Recall $v(1), \ldots, v(n)$ are all of the available denominations.)
- Among all of these optimal solutions (which we find in the table we are constructing recursively!) we pick one which uses the fewest number of coins, say this is opt(i-v(m)) for some m, 1 < m < n.
- We obtain an optimal solution opt(i) for amount i by adding to opt(i-v(m)) one coin of denomination v(m).
- Why does this produce an optimal solution for amount $i \leq C$?
- Consider an optimal solution for amount $i \leq C$; and say such solution includes at least one coin of denomination v(m) for some $1 \leq m \leq n$. But then removing such a coin must produce an optimal solution for the amount i v(m) again by our cut-and-paste argument.
- However, we do not know which coins the optimal solution includes, so we try all the available coins and then pick m for which the optimal solution for amount i-v(m) uses the fewest number of coins.

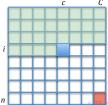
- It is enough to store in the i^{th} slot of the table such m and opt(i) because this allows us to reconstruct the optimal solution by looking at m_1 stored in the i^{th} slot, then look at m_2 stored in the slot $i v(m_1)$, then look at m_2 stored in the slot $i v(m_1) v(m_2)$, etc.
- Finally, after C many such steps we obtain opt(C) which is the solution we need.
- Time complexity of our algorithm is nC.
- Note: Our algorithm is NOT a polynomial time algorithm in the **length** of the input, because the length of a representation of C is only $\log C$, while the running time is nC.
- But this is the best what we can do...

Integer Knapsack Problem (Duplicate Items Allowed) You have n types of items; all items of kind i are identical and of weight w_i and value v_i . You also have a knapsack of capacity C. Choose a combination of available items which all fit in the knapsack and whose value is as large as possible. You can take any number of items of each kind.

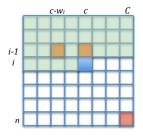
- ullet Solution: DP recursion on the capacity C of the knapsack.
- We build a table of optimal solutions for all knapsacks of capacities $i \leq C$.
- Assume we have solved the problem for all knapsacks of capacities j < i.
- We now look at optimal solutions $opt(i-w_m)$ for all knapsacks of capacities $i-w_m$ for all $1 \le m \le n$;
- Chose the one for which $opt(i-w_m) + v_m$ is the largest;
- Add to such optimal solution for the knapsack of size $i w_m$ item m to obtain a packing of a knapsack of size i of the highest possible value.

- Thus, $opt(i) = \max\{opt(i w_m) + v_m : 1 \le m \le n\}.$
- After C many steps we obtain opt(C) which is what we need.
- Again, our algorithm is **NOT** polynomial in the **length** of the input.

- Integer Knapsack Problem (Duplicate Items NOT Allowed) You have n items (some of which can be identical); item I_i is of weight w_i and value v_i . You also have a knapsack of capacity C. Choose a combination of available items which all fit in the knapsack and whose value is as large as possible.
- This is an example of a "2D" recursion; we will be filling a table of size $n \times C$, row by row; subproblems P(i,c) for all $i \le n$ and $c \le C$ will be of the form:
 - chose from items I_1, I_2, \ldots, I_i a subset which fits in a knapsack of capacity c and is of the largest possible total value.
- Fix now $i \leq n$ and $c \leq C$ and assume we have solved the subproblems for:
 - **1** all j < i and all knapsacks of capacities from 1 to C;
 - 2 for i we have solved the problem for all capacities d < c.



- we now have two options: either we take item I_i or we do not;
- so we look look at optimal solutions $opt(i-1, c-w_i)$ and opt(i-1, c);



- if $opt(i-1, c-w_i) + v_i > opt(i-1, c)$ then $opt(i, c) = opt(i-1, c-w_i) + v_i;$ else opt(i, c) = opt(i-1, c).
- Final solution will be given by opt(n, C).

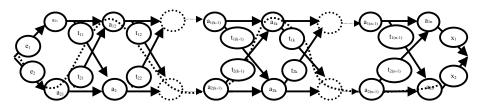
- Balanced Partition You have a set of n integers. Partition these integers into two subsets such that you minimise $|S_1 S_2|$, where S_1 and S_2 denote the sums of the elements in each of the two subsets.
- Solution: Let S be the total sum of all integers in the set; consider the Knapsack problem (with duplicate items not allowed) with the knapsack of size S/2 and with each integer x_i of both size and value equal to x_i .
- Claim: the best packing of such knapsack produces optimally balanced partition, with S_1 being all the integers in the knapsack and S_2 all the integers left out of the knapsack.
- Why? Since $S = S_1 + S_2$ we obtain

$$\frac{S}{2} - S_1 = \frac{S_1 + S_2}{2} - S_1 = \frac{S_2 - S_1}{2}$$

i.e.
$$S_2 - S_1 = 2(S/2 - S_1)$$
.

- Thus, minimising $S/2 S_1$ will minimise $S_2 S_1$.
- So, all we have to do is find a subset of these numbers with the largest possible total sum which fits inside a knapsack of size S/2.

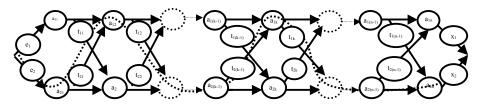
Dynamic Programming: Assembly line scheduling



Instance: Two assembly lines with workstations for n jobs.

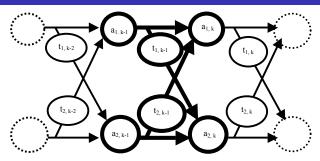
- On the first assembly line the k^{th} job takes $a_{1,k}$ $(1 \le k \le n)$ units of time to complete; on the second assembly line the same job takes $a_{2,k}$ units of time.
- To move the product from station k-1 on the first assembly line to station k on the second line it takes $t_{1,k-1}$ units of time.
- Likewise, to move the product from station k-1 on the second assembly line to station k on the first assembly line it takes $t_{2,k-1}$ units of time.

Dynamic Programming: Assembly line scheduling



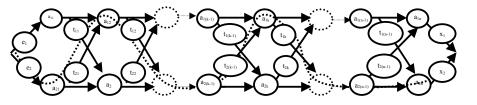
- To bring an unfinished product to the first assembly line it takes e_1 units of time.
- ullet To bring an unfinished product to the second assembly line it takes e_2 units of time.
- To get a finished product from the first assembly line to the warehouse it takes x_1 units of time;
- ullet To get a finished product from the second assembly line to the warehouse it takes x_2 units.
- Task: Find a fastest way to assemble a product using both lines as necessary.

Dynamic Programming: Assembly line scheduling



- For each $k \le n$, we solve subproblems P(1,k) and P(2,k) by a **simultaneous** recursion on k:
- P(1,k): find the minimal amount of time m(1,k) needed to finish the first k jobs, such the k^{th} job is finished on the k^{th} workstation on the **first** assembly line;
- P(2,k): find the minimal amount of time m(2,k) needed to finish the first k jobs, such the k^{th} job is finished on the k^{th} workstation on the **second** assembly line.

Dynamic Programming



- We solve P(1, k) and P(2, k) by a simultaneous recursion:
- Initially $m(1,1) = e_1 + a_{1,1}$ and $m(2,1) = e_2 + a_{2,1}$.
- Recursion:

$$\begin{split} m(1,k) &= \min\{m(1,k-1) + a_{1,k}, \quad m(2,k-1) + t_{2,k-1} + a_{1,k}\} \\ m(2,k) &= \min\{m(2,k-1) + a_{2,k}, \quad m(1,k-1) + t_{1,k-1} + a_{2,k}\} \end{split}$$

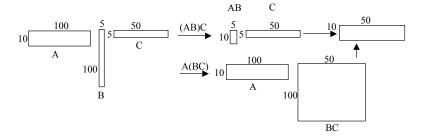
• Finally, after obtaining m(1, n) and m(2, n) we choose

$$opt = min\{m(1, n) + x_1, m(2, n) + x_2\}.$$

• This problem is important because it has the same design logic as the Viterbi algorithm, an extremely important algorithm for many fields such as speech recognition, decoding convolutional codes in telecommunications etc, covered in COMP4121 (Advanced Algorithms).

- For any three matrices of compatible sizes we have A(BC) = (AB)C.
- However, the number of real number multiplications needed to perform in order to obtain the matrix product can be very different:

$$A = 10 \times 100;$$
 $B = 100 \times 5;$ $C = 5 \times 50$



- To evaluate (AB)C we need $(10 \times 5) \times 100 + (10 \times 50) \times 5 = 5000 + 2500 = 7500$ multiplications;
- To evaluate A(BC) we need $(100 \times 50) \times 5 + (10 \times 50) \times 100 = 25000 + 50000 = 75000$ multiplications!

- Problem Instance: A sequence of matrices $A_1A_2...A_n$;
- Task: Group them in such a way as to minimise the total number of multiplications needed to find the product matrix.
- \bullet The total number of different distributions of brackets is equal to the number of binary trees with n leaves.
- The total number of different distributions of brackets satisfies the following recursion (why?):

$$T(n) = \sum_{i=1}^{n-1} T(i)T(n-i)$$

- One can show that the solution satisfies $T(n) = \Omega(2^n)$.
- Thus, we cannot do an exhaustive search for the optimal placement of the brackets.

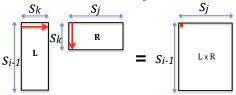


- Problem Instance: A sequence of matrices $A_1A_2...A_n$;
- Task: Group them in such a way as to minimise the total number of multiplications needed to find the product matrix.
- The subproblems P(i, j) to be considered are:
 - "group matrices $A_iA_{i+1}...A_{j-1}A_j$ in such a way as to minimise the total number of multiplications needed to find the product matrix".
- Note: this looks like it is a case of a "2D recursion, but we can actually do it with a simple "linear" recursion.
- We group such subproblems by the value of j-i and perform a recursion on the value of j-i.
- At each recursive step m we solve all subproblems P(i,j) for which j-i=m.



- Let m(i,j) denote the minimal number of multiplications needed to compute the product $A_i A_{i+1} ... A_{j-1} A_j$; let also the size of matrix A_i be $s_{i-1} \times s_i$.
- Recursion: We examine all possible ways to place the principal (outermost) multiplication, splitting the chain into the product $(A_i \dots A_k) \cdot (A_{k+1} \dots A_j)$.
- Note that both k-i < j-i and j-(k+1) < j-i; thus we have the solutions of the subproblems P(i,k) and P(k+1,j) already computed and stored in slots k-i and j-(k+1), respectively, which precede slot j-i we are presently filling.
- Note also that the matrix product $A_i \dots A_k$ is a $s_{i-1} \times s_k$ matrix L and $A_{k+1} \dots A_j$ is a $s_k \times s_j$ matrix R.

• To multiply an $s_{i-1} \times s_k$ matrix L and an $s_k \times s_j$ matrix R it takes $s_{i-1}s_ks_j$ many multiplications: S_k S_i S_j



Total number of multiplications: S_{i-1} S_j S_k

• The recursion:

$$m(i,j) = \min\{m(i,k) + m(k+1,j) + s_{i-1}s_js_k : i \le k \le j-1\}$$

- Note that the recursion step is a brute force search but the whole algorithm is not, because all the subproblems are solved only once, and there are only $O(n^2)$ many such subproblems.
- k for which the minimum in the recursive definition of m(i,j) is achieved can be stored to retrieve the optimal placement of brackets for the whole chain $A_1 \ldots A_n$.
- Thus, in the m^{th} slot of the table we are constructing we store all pairs (m(i,j),k) for which j-i=m.

- Assume we want to compare how similar two sequences of symbols S and S^* are.
- Example: how similar are the genetic codes of two viruses.
- This can tell us if one is just a genetic mutation of the other.
- A sequence s is a **subsequence** of another sequence S if s can be obtained by deleting some of the symbols of S (while preserving the order of the remaining symbols).
- Given two sequences S and S^* a sequence s is a **Longest** Common Subsequence of S, S^* if s is a common subsequence of both S and S^* and is of maximal possible length.

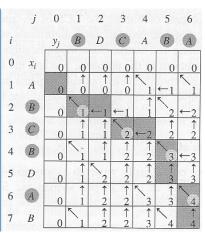
- Instance: Two sequences $S = \langle a_1, a_2, \dots a_n \rangle$ and $S^* = \langle b_1, b_2, \dots, b_m \rangle$.
- Task: Find a longest common subsequence of S, S^* .
- We first find the length of the longest common subsequence of S, S^* .
- "2D recursion": for all $1 \le i \le n$ and all $1 \le j \le m$ let c[i,j] be the length of the longest common subsequence of the truncated sequences $S_i = \langle a_1, a_2, \dots a_i \rangle$ and $S_j^* = \langle b_1, b_2, \dots, b_j \rangle$.
- Recursion: we fill the table row by row, so the ordering of subproblems is the lexicographical ordering;

$$c[i,j] = \begin{cases} 0, & \text{if } i = 0 \text{ or } j = 0; \\ c[i-1,j-1] + 1 & \text{if } i,j > 0 \text{ and } a_i = b_j; \\ \max\{c[i-1,j], c[i,j-1]\} & \text{if } i,j > 0 \text{ and } a_i \neq b_j. \end{cases}$$



Retrieving a longest common subsequence:

```
LCS-LENGTH(X, Y)
      m \leftarrow length[X]
     n \leftarrow length[Y]
      for i \leftarrow 1 to m
            do c[i, 0] \leftarrow 0
      for j \leftarrow 0 to n
            do c[0, j] \leftarrow 0
      for i \leftarrow 1 to m
  8
            do for j \leftarrow 1 to n
                      do if x_i = y_i
                              then c[i, j] \leftarrow c[i - 1, j - 1] + 1
                                    b[i, j] \leftarrow "\"
                              else if c[i-1, j] > c[i, j-1]
                                       then c[i, j] \leftarrow c[i-1, j]
                                              b[i, j] \leftarrow "\uparrow"
15
                                       else c[i, j] \leftarrow c[i, j-1]
                                              b[i, i] \leftarrow "\leftarrow"
      return c and b
```



- What if we have to find a longest common subsequence of three sequences S_1, S_2, S_3 ?
- Can we do $LCS(LCS(S_1, S_2), S_3)$?
- Not necessarily! Consider

$$S_1 = ABCDEGG$$
 $LCS(S_1, S_2) = ABEG$
 $S_2 = ACBEEFG$ $LCSS_2, S_3) = ACEF$
 $S_3 = ACCEDGF$ $LCS(S_1, S_3) = ACDG$

$$\begin{split} & \operatorname{LCS}(\operatorname{LCS}(S_1, S_2), S_3) = \operatorname{LCS}(ABEG, ACCEDGF) = AEG \\ & \operatorname{LCS}(\operatorname{LCS}(S_2, S_3), S_1) = \operatorname{LCS}(ACEF, ABCDEGG) = ACE \\ & \operatorname{LCS}(\operatorname{LCS}(S_1, S_3), S_2) = \operatorname{LCS}(ACDG, ACBEEFG) = ACG \end{split}$$

But

$$LCS(S_1, S_2, S_3) = ACEG$$

• So how would you design an algorithm which computes correctly $LCS(S_1, S_2, S_3)$?



- Instance: Three sequences $S = \langle a_1, a_2, \dots a_n \rangle$, $S^* = \langle b_1, b_2, \dots, b_m \rangle$ and $S^{**} = \langle c_1, c_2, \dots, c_k \rangle$.
- Task: Find a longest common subsequence of S, S^*, S^{**} .
- ullet We again first find the length of the longest common subsequence of S, S^*, S^{**} .
- for all $1 \le i \le n$, all $1 \le j \le m$ and all $l \le k$, let d[i, j, l] be the length of the longest common subsequence of the truncated sequences $S_i = \langle a_1, a_2, \dots a_i \rangle$, $S_j^* = \langle b_1, b_2, \dots, b_j \rangle$ and $S_l^* = \langle c_1, c_2, \dots, c_l \rangle$.
- Recursion:

$$d[i,j,l] = \begin{cases} 0, & \text{if } i=0 \text{ or } j=0 \text{ or } l=0; \\ d[i-1,j-1,l-1]+1 & \text{if } i,j,l>0 \text{ and } a_i=b_j=c_l; \\ \max\{d[i-1,j,l],d[i,j-1,l],d[i,j,l-1]\} & \text{otherwise.} \end{cases}$$



Dynamic Programming: Shortest Common Supersequence

- Instance: Two sequences $s = \langle a_1, a_2, \dots a_n \rangle$ and $s^* = \langle b_1, b_2, \dots, b_m \rangle$.
- Task: Find a shortest common super-sequence S of s, s^* , i.e., the shortest possible sequence S such that both s and s^* are subsequences of S.
- Solution: Find the longest common subsequence $LCS(s, s^*)$ of s and s^* and then add differing elements of the two sequences at the right places, in any order; for example:

$$s = a\mathbf{b}a\mathbf{c}ada$$

 $s^* = x\mathbf{b}y\mathbf{c}az\mathbf{d}$
 $LCS(s, s^*) = \mathbf{b}cad$
shortest super-sequence $S = ax\mathbf{b}ya\mathbf{c}az\mathbf{d}a$

Dynamic Programming: Edit Distance

- Edit Distance Given two text strings A of length n and B of length m, you want to transform A into B. You are allowed to insert a character, delete a character and to replace a character with another one. An insertion costs c_i , a deletion costs c_d and a replacement costs c_r .
- Task: find the lowest total cost transformation of A into B.
- Note: if all operations have a unit cost, then you are looking for the minimal number of such operations required to transform A into B; this number is called *the edit distance* between A and B.
- If the sequences are sequences of DNA bases and the costs reflect the probabilities of the corresponding mutations, then the minimal cost represents how closely related the two sequences are.
- Subproblems: Let C(i,j) be the minimum cost of transforming the sequence A[1..i] into the sequence B[1..j] for all $i \leq n$ and all $j \leq m$.



Dynamic Programming: Edit Distance

- Subproblems P(i,j): Find the minimum cost C(i,j) of transforming the sequence A[1..i] into the sequence B[1..j] for all $i \leq n$ and all $j \leq m$.
- Recursion: we again fill the table of solutions C(i, j) for subproblems P(i, j) row by row (why is this OK?):

$$C(i,j) = \min \begin{cases} c_d + C(i-1,j) \\ C(i,j-1) + c_i \\ \begin{cases} C(i-1,j-1) & \text{if } A[i] = B[j] \\ C(i-1,j-1) + c_r & \text{if } A[i] \neq B[j] \end{cases} \end{cases}$$

- cost $c_d + C(i-1,j)$ corresponds to the option if you first delete A[i] and then recursively transform A[1..i-1] into B[1..j];
- cost $C(i, j-1) + c_i$ corresponds to the option if you first transform A[1..i] to B[1..j-1] and then append B[j] at the end;
- the third option corresponds to first transforming A[1..i-1] to B[1..j-1] and
 - if A[i] is already equal to B[j] do nothing, thus incurring a cost of only C(i-1, j-1);
 - ② if A[i] is not equal to B[j] replace A[i] by B[j] with a total cost of $C(i-1,j-1)+c_r$.

Dynamic Programming: Maximizing an expression

• Instance: a sequence of numbers with operations $+, -, \times$ in between, for example

$$1 + 2 - 3 \times 6 - 1 - 2 \times 3 - 5 \times 7 + 2 - 8 \times 9$$

- Task: Place brackets in a way that the resulting expression has the largest possible value.
- What will be the subproblems?
- Maybe for a subsequence of numbers A[i..j] place the brackets so that the resulting expression is maximised?
- maybe we could consider which the principal operations should be, i.e. $A[i..k] \odot A[k+1..j]$. Here \odot is whatever operation is between A[k] and A[k+1].
- But when would such expression be maximised if there could be both positive and negative values for A[i..k] and A[k+1..j] depending on the placement of brackets??
- Maybe we should look for placements of brackets not only for the maximal value but also for the minimal value!
- Exercise: write the exact recursion for this problem.



Dynamic Programming: Turtle Tower

- Instance: You are given n turtles, and for each turtle you are given its weight and its strength. The strength of a turtle is the maximal weight you can put on it without cracking its shell.
- Task: Find the largest possible number of turtles which you can stack one on top of the other, without cracking any turtle.
- **Hint:** Order turtles in an increasing order of the sum of their weight and their strength, and proceed by recursion.
- You can find a solution to this problem and of another interesting problem on the class website (class resources, file "More Dynamic Programming").

Dynamic Programming: Bellman Ford algorithm

- $\bullet\,$ One of the earliest use of Dynamic Programming (1950's) invented by Bellman.
- Instance: A directed weighted graph G = (V, E) with weights which can be negative, but without cycles of negative total weight and a vertex $s \in V$.
- Goal: Find the shortest path from vertex s to every other vertex t.
- Solution: Since there are no negative weight cycles, the shortest path cannot contain cycles, because a cycle can be excised to produce a shorter path.
- Thus, every shortest path can have at most |V|-1 edges.
- Subproblems: For every $v \in V$ and every i, $(1 \le i \le n-1)$, let opt(i, v) be the length of a shortest path from s to v which contains at most i edges.
- Our goal is to find for every vertex $t \in G$ the value of $\operatorname{opt}(n-1,t)$ and the path which achieves such a length.
- Note that if the shortest path from a vertex v to t is $(v, p_1, p_2, \ldots, p_k, t)$ then $(p_1, p_2, \ldots, p_k, t)$ must be the shortest path from p_1 to t, and $(v, p_1, p_2, \ldots, p_k)$ must also be the shortest path from v to p_k .

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Dynamic Programming: Bellman Ford algorithm

• Let us denote the length of the shortest path from s to v among all paths which contain at most i edges by $\operatorname{opt}(i,v)$, and let $\operatorname{pred}(i,v)$ be the immediate predecessor of vertex v on such shortest path.

• Recursion:

$$\begin{aligned} & \operatorname{opt}(i,v) = \min(\operatorname{opt}(i-1,v), \min_{p \in V} \{\operatorname{opt}(i-1,p) + \operatorname{w}(e(p,v))\}; \\ & \operatorname{pred}(i,v) = \begin{cases} \operatorname{pred}(i-1,v) & \text{if } \min_{p \in V} \{\operatorname{opt}(i-1,p) + \operatorname{w}(e(p,v))\} \geq \operatorname{pred}(i-1,v) \\ \operatorname{arg\,min}_{p \in V} \{\operatorname{opt}(i-1,p) + \operatorname{w}(e(p,v))\} & \text{otherwise} \end{cases} \end{aligned}$$

(here w(e(p, v)) is the weight of the edge e(p, v) from vertex p to vertex v.)

- Computation $\operatorname{opt}(i,v)$ runs in time $O(|V| \times |E|)$, because $i \leq |V| 1$ and for each v, min is taken over all edges e(p,v) incident to v; thus in each round all edges are inspected.
- ullet Algorithm produces shortest paths from s to every other vertex in the graph.
- The method employed is sometimes called "relaxation", because we progressively relax the additional constraint on how many edges the shortest paths can contain.

Dynamic Programming: Floyd Warshall algorithm

- Let again G = (V, E) be a directed weighted graph where $V = \{v_1, v_2, \dots, v_n\}$ and where weights $w(e(v_p, v_q))$ of edges $e(v_p, v_q)$ can be negative, but there are no negative weight cycles.
- We can use a somewhat similar idea to obtain the shortest paths from **every** vertex v_p to **every** vertex v_q (including back to v_p).
- Let $\operatorname{opt}(k, v_p, v_q)$ be the length of the shortest path from a vertex v_p to a vertex v_q such that all intermediate vertices are among vertices $\{v_1, v_2, \ldots, v_k\}, (1 \leq k \leq n).$
- Then

$$opt(k, v_p, v_q) = \min\{opt(k - 1, v_p, v_q), opt(k - 1, v_p, v_k) + opt(k - 1, v_k, v_q)\}$$

- Thus, we gradually **relax** the constraint that the intermediary vertices have to belong to $\{v_1, v_2, \dots, v_k\}$.
- Algorithm runs in time $|V|^3$.



Dynamic Programming: relaxation

• Compute the number of partitions of a positive integer n. That is to say the number of distinct **sets** of positive integers $\{n_1, \ldots, n_k\}$ which sum up to n, i.e., such that $n_1 + \ldots + n_k = n$.

Hint: Let $\operatorname{nump}(i,j)$ denotes the number of partitions $\{j_1,\ldots,j_p\}$ of j, i.e., the number of sets such that $j_1+\ldots+j_p=j$ which, in addition, have the property that every element j_q of each partition satisfies $j_q \leq i$. We are looking for $\operatorname{nump}(n,n)$ but the recursion is based on relaxation of the allowed size i of the parts of j for all $i,j\leq n$. To get a recursive definition of $\operatorname{nump}(i,j)$ distinguish the case of partitions where all components are $\leq i-1$ and the case where at least one component is of size i.