



Chapter 3

Dynamic Programming

Algorithm Theory
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Dynamic Programming



„Memoization“ for increasing the efficiency of a recursive solution:

- Only the *first time* a sub-problem is encountered, its **solution is computed** and then stored in a table. Each subsequent time that the subproblem is encountered, the value stored in the table is simply looked up and returned
(without repeated computation!).
- **Computing the solution**: For each sub-problem, store how the value is obtained (according to which recursive rule).

Dynamic Programming

Dynamic programming / memoization can be applied if

- **Optimal solution** contains **optimal solutions to sub-problems**
(recursive structure)
- Number of sub-problems that need to be considered is small

Knapsack

- n items $1, \dots, n$, each item has **weight w_i** and **value v_i**
- Knapsack (bag) of capacity W
- Goal: pack items into knapsack such that **total weight** is at most W and **total value is maximized**:

$$\begin{aligned} & \max \sum_{i \in S} v_i \\ & \text{s. t. } \underline{S} \subseteq \{1, \dots, n\} \text{ and } \sum_{i \in S} w_i \leq W \end{aligned}$$

- E.g.: jobs of length w_i and value v_i , server available for W time units, try to execute a set of jobs that maximizes the total value

Recursive Structure?

- Optimal solution: \mathcal{O} *cases : $n \in \mathcal{O}, n \notin \mathcal{O}$*
- If $n \notin \mathcal{O}$: $\text{OPT}(n) = \text{OPT}(n - 1)$
- What if $n \in \mathcal{O}$?
 - Taking n gives value v_n
 - But, n also occupies space w_n in the bag (knapsack)
 - There is space for $W - w_n$ total weight left!

$\text{OPT}(n) = \overset{v_n}{\cancel{w_n}} + \text{optimal solution with first } n - 1 \text{ items}$
and knapsack of capacity $W - w_n$

A More Complicated Recursion

$OPT(k, x)$: value of optimal solution with items 1, ..., k and knapsack of capacity x

Recursion:

$$OPT(k, x) = \max \left\{ \underbrace{OPT(k-1, x)}_{\substack{\text{opt. solution} \\ \text{does not contain} \\ \text{item k}}}, v_k + \underbrace{OPT(k-1, x-w_k)}_{\substack{\text{remaining} \\ \text{capacity} \\ \text{when adding} \\ \text{item k}}} \right\}$$

only makes sense if $x - w_k \geq 0$

items $1 \dots k$ capacity of knaps.

initialisation

$$\underline{OPT(0, x) = 0}$$

$$\underline{OPT(k, 0) = 0}$$

exp. / ∞ many options for 2nd parameter

assumption: weights are integers

\Rightarrow only $W+1$ options for 2nd parameter

Dynamic Programming Algorithm

Set up table for all possible $OPT(k, x)$ -values

- Assume that all weights w_i are integers!

	0	1	2	3	...	W
1						
2						
3				<u>$OPT(3,5)$</u>		
...						
n						

Handwritten annotations:
 - A vertical arrow on the left side of the table is labeled "items".
 - A horizontal arrow points from the "3" in the row index to the cell containing $OPT(3,5)$.
 - A vertical arrow points from the "5" in the column index to the same cell.

Row i , column j :
 $OPT(i, j)$

Example

- 1 2 3 4 5 6 7 8
- 8 items: (3,2), (2,4), (4,1), (5,6), (3,3), (4,3), (5,4), (6,6)
 Knapsack capacity: 12
- weight value

- $OPT(k, x) = \max\{OPT(k-1, x), OPT(k-1, x-w_k) + v_k\}$

	1	2	3	4	5	6	7	8	9	10	11	12
1	0	0	2	2	2	2	2	2	2	2	2	2
2	0	4	4	4	6	6	6	6	6	6	6	6
3	0	4	4	4	6	6	6	6	7	7	7	7
4												
5												
6												
7												
8												

← only item 1

Running Time of Knapsack Algorithm

- **Size of table:** $O(n \cdot W)$
- Time per table entry: $O(1)$ → **overall time:** $O(nW)$
- Computing solution (set of items to pick):
Follow $\leq n$ arrows → $O(n)$ time (after filling table)
- Note: Time depends on W → can be exponential in n ...
- And it is problematic if weights are not integers.

another special case : values are integers

general : problem is NP-hard

but: we can get arbitrarily good solutions efficiently

String Matching Problems

Edit distance $D(A, B)$ (between strings A and B):

m a - t h e m - - a t i c i a n
m u l t i p l i c a t i o - - n

Approximate string matching:

For a given text T , a pattern P and a distance d , find all
substrings P' of T with $D(P, P') \leq d$.

Sequence alignment:

Find optimal alignments of DNA / RNA / ... sequences.

G A G C A - C T T G G A T T C T C G G
- - - C A C G T G G - A - A C T - - -

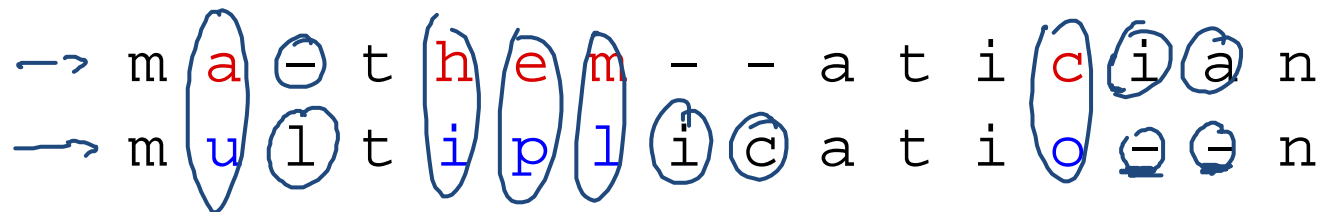
Edit Distance

Given: Two strings $A = a_1 a_2 \dots a_m$ and $B = b_1 b_2 \dots b_n$

Goal: Determine the minimum number $D(A, B)$ of edit operations required to transform A into B

Edit operations:

- a) **Replace** a character from string A by a character from B
- b) **Delete** a character from string A
- c) **Insert** a character from string B into A



Edit Distance – Cost Model

- Cost for **replacing** character a by b : $c(a, b) \geq 0$
- Capture insert, delete by allowing $a = \varepsilon$ or $b = \varepsilon$:
 - Cost for **deleting** character a : $c(a, \varepsilon)$
 - Cost for **inserting** character b : $c(\varepsilon, b)$

- **Triangle inequality:**

$$\underline{c(a, c)} \leq \underline{c(a, b)} + \underline{c(b, c)}$$

→ each character is changed at most once!

- **Unit cost model:** $c(a, b) = \begin{cases} 1, & \text{if } a \neq b \\ 0, & \text{if } a = b \end{cases}$ } counting # of edit operations

Recursive Structure

- Optimal “alignment” of strings (unit cost model)

bbcadfagikccm and abbagflrgikacc:

-	b	b	c	a	g	f	a		-	g	i	k	-	c	c	m
a	b	b	-	a	d	f	l	r	g	i	k	a	c	c	-	

- Consists of optimal “alignments” of sub-strings, e.g.:

-bbcagfa abb-adfl	and	-gik-ccm rgikacc-
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- Edit distance between $A_{1,m} = a_1 \dots a_m$ and $B_{1,n} = b_1 \dots b_n$:

$$\underline{D(A, B)} = \min_{\underline{k, \ell}} \{ \underline{D(A_{1,k}, B_{1,\ell})} + \underline{D(A_{k+1,m}, B_{\ell+1,n})} \}$$

Computation of the Edit Distance

string $A = a_1, \dots, a_m$, $B = b_1, \dots, b_n$

Let $A_k := a_1 \dots a_k$, $B_\ell := b_1 \dots b_\ell$, and

$$\underline{\underline{D_{k,\ell}}} := \underline{\underline{D(A_k, B_\ell)}}$$

