

17. Dynamic Programming II

Editing Distance, Bellman-Ford Algorithm
[Cormen et al, Kap. 24.1]]

Minimal Editing Distance

Editing distance of two sequences $A_n = (a_1, \dots, a_n)$, $B_m = (b_1, \dots, b_m)$.

Editing operations:

- Insertion of a character
- Deletion of a character
- Replacement of a character

Question: how many editing operations at least required in order to transform string A into string B .

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Minimal Editing Distance

Wanted: cheapest character-wise transformation $A_n \rightarrow B_m$ with costs

operation	Levenshtein	LCS ²⁴	general
Insert c	1	1	$\text{ins}(c)$
Delete c	1	1	$\text{del}(c)$
Replace $c \rightarrow c'$	$\mathbb{1}(c \neq c')$	$\infty \cdot \mathbb{1}(c \neq c')$	$\text{repl}(c, c')$

Beispiel

T I G E R
Z I E G E

T I _ G E R
Z I E G E _

T → Z +E -R
Z → T -E +R

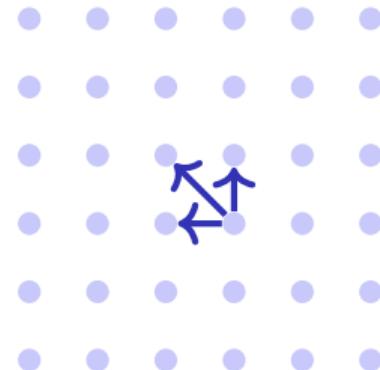
²⁴Longest common subsequence – A special case of an editing problem

0. $E(n, m)$ = minimum number edit operations (ED cost) $a_{1..n} \rightarrow b_{1..m}$
1. Subproblems $E(i, j)$ = ED von $a_{1..i}, b_{1..j}$. #SP = $n \cdot m$
2. Guess Costs $\Theta(1)$
 - $a_{1..i} \rightarrow a_{1..i-1}$ (delete)
 - $a_{1..i} \rightarrow a_{1..i}b_j$ (insert)
 - $a_{1..i} \rightarrow a_{1..i-1}b_j$ (replace)

3. Rekursion

$$E(i, j) = \min \begin{cases} \text{del}(a_i) + E(i - 1, j), \\ \text{ins}(b_j) + E(i, j - 1), \\ \text{repl}(a_i, b_j) + E(i - 1, j - 1) \end{cases}$$

4. Dependencies



⇒ Computation from left top to bottom right. Row- or column-wise.

5. Solution in $E(n, m)$

Example (Levenshtein Distance)

$$E[i, j] \leftarrow \min \{E[i - 1, j] + 1, E[i, j - 1] + 1, E[i - 1, j - 1] + \mathbb{1}(a_i \neq b_j)\}$$

	\emptyset	Z	I	E	G	E
\emptyset	0	1	2	3	4	5
T	1	1	2	3	4	5
I	2	2	1	2	3	4
G	3	3	2	2	2	3
E	4	4	3	2	3	2
R	5	5	4	3	3	3

Editing steps: from bottom right to top left, following the recursion.
Bottom-Up description of the algorithm: exercise

Bottom-Up DP algorithm ED

Dimension of the table? Semantics?

1.

Bottom-Up DP algorithm ED

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1. Table $E[0, \dots, m][0, \dots, n]$. $E[i, j]$: minimal edit distance of the strings (a_1, \dots, a_i) and (b_1, \dots, b_j)

Bottom-Up DP algorithm ED

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Computation of an entry

- 2.

Bottom-Up DP algorithm ED

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1. Table $E[0, \dots, m][0, \dots, n]$. $E[i, j]$: minimal edit distance of the strings (a_1, \dots, a_i) and (b_1, \dots, b_j)

Computation of an entry

2. $E[0, i] \leftarrow i \ \forall 0 \leq i \leq m, E[j, 0] \leftarrow i \ \forall 0 \leq j \leq n$. Computation of $E[i, j]$ otherwise via $E[i, j] = \min\{\text{del}(a_i) + E(i-1, j), \text{ins}(b_j) + E(i, j-1), \text{repl}(a_i, b_j) + E(i-1, j-1)\}$

Bottom-Up DP algorithm ED

3.

Computation order

Bottom-Up DP algorithm ED

Computation order

3.

Rows increasing and within columns increasing (or the other way round).

Bottom-Up DP algorithm ED

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Rows increasing and within columns increasing (or the other way round).
4. Reconstruction of a solution?

Bottom-Up DP algorithm ED

Computation order

3.

Rows increasing and within columns increasing (or the other way round).

Reconstruction of a solution?

4. Start with $j = m, i = n$. If $E[i, j] = \text{repl}(a_i, b_j) + E(i - 1, j - 1)$ then output $a_i \rightarrow b_j$ and continue with $(j, i) \leftarrow (j - 1, i - 1)$; otherwise, if $E[i, j] = \text{del}(a_i) + E(i - 1, j)$ output $\text{del}(a_i)$ and continue with $j \leftarrow j - 1$ otherwise, if $E[i, j] = \text{ins}(b_j) + E(i, j - 1)$, continue with $i \leftarrow i - 1$. Terminate for $i = 0$ and $j = 0$.

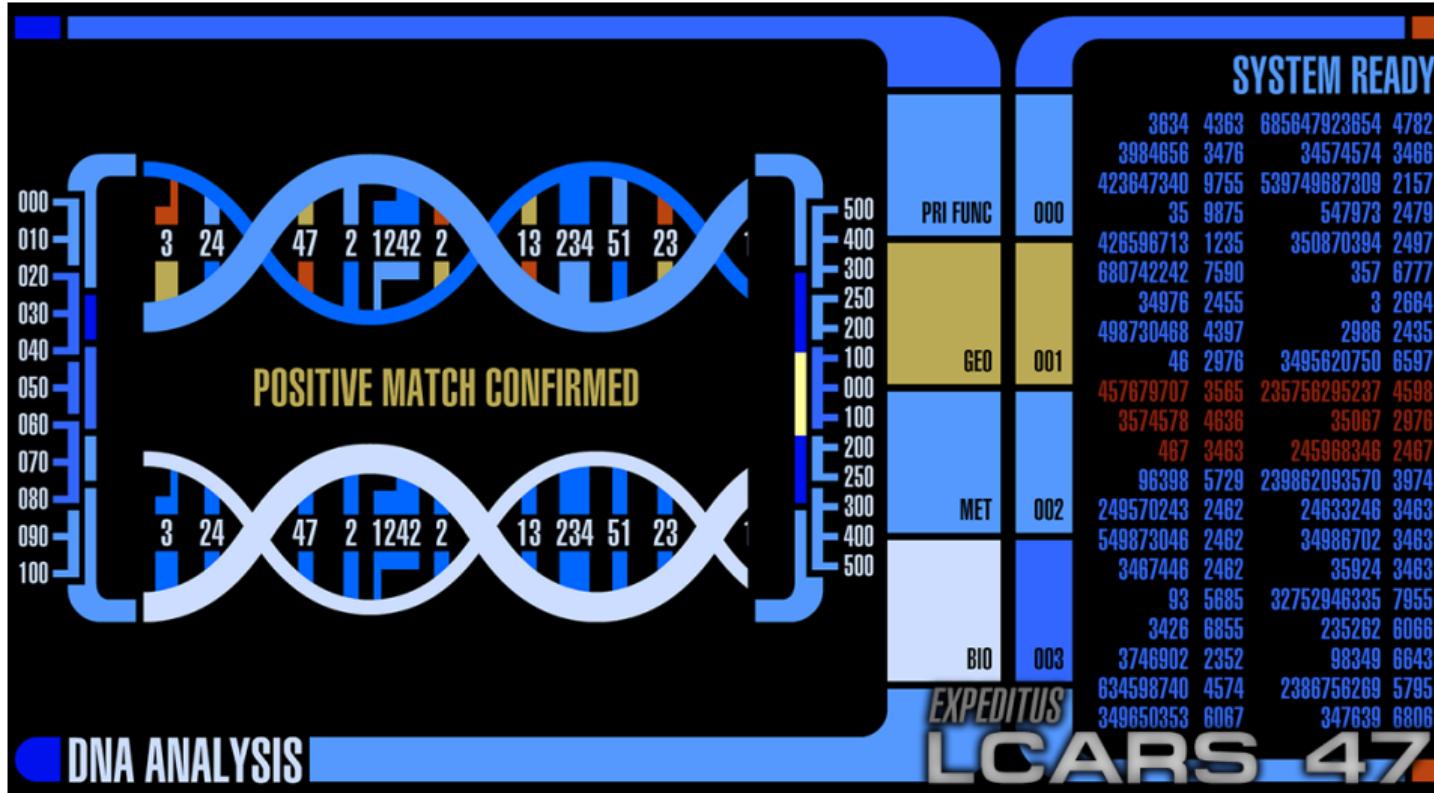
Analysis ED

- Number table entries: $(m + 1) \cdot (n + 1)$.
- Constant number of assignments and comparisons each. Number steps: $\mathcal{O}(mn)$
- Determination of solution: decrease i or j . Maximally $\mathcal{O}(n + m)$ steps.

Runtime overall:

$$\mathcal{O}(mn).$$

DNA - Comparison (Star Trek)



DNA - Comparison

- DNA consists of sequences of four different nucleotides **A**dénine
Guanine **T**hymine **C**ytosine
- DNA sequences (genes) thus can be described with strings of A, G, T and C.
- Possible comparison of two genes: determine the **longest common subsequence**

The longest common subsequence problem is a special case of the minimal edit distance problem.

Reminder: Shortest Path Algorithm

1. Initialise d_s and π_s : $d_s[v] = \infty$, $\pi_s[v] = \text{null}$ for each $v \in V$
2. Set $d_s[s] \leftarrow 0$
3. Choose an edge $(u, v) \in E$

Relaxiere (u, v) :

```
if  $d_s[v] > d_s[u] + c(u, v)$  then  
     $d_s[v] \leftarrow d_s[u] + c(u, v)$   
     $\pi_s[v] \leftarrow u$ 
```

4. Repeat 3 until nothing can be relaxed any more.
 $(\text{until } d_s[v] \leq d_s[u] + c(u, v) \quad \forall (u, v) \in E)$

Dynamic Programming Approach (Bellman)

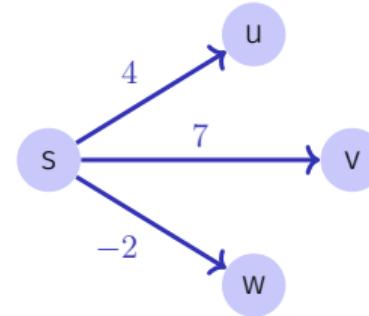
Induction over number of edges $d_s[i, v]$: Shortest path from s to v via maximally i edges.

$$d_s[i, v] = \min\{d_s[i - 1, v], \min_{(u,v) \in E} (d_s[i - 1, u] + c(u, v))\}$$

$$d_s[0, s] = 0, d_s[0, v] = \infty \quad \forall v \neq s.$$

Dynamic Programming Approach (Bellman)

	s	\dots	v	\dots	w
0	0	∞	∞	∞	∞
1	0	∞	7	∞	-2
\vdots	\vdots	\vdots	\vdots	\vdots	\vdots
$n - 1$	0	\dots	\dots	\dots	\dots



Algorithm: Iterate over last row until the relaxation steps do not provide any further changes, maximally $n - 1$ iterations. If still changes, then there is no shortest path.

Algorithm Bellman-Ford(G, s)

Input: Graph $G = (V, E, c)$, starting point $s \in V$

Output: If return value true, minimal weights d for all shortest paths from s , otherwise no shortest path.

```
foreach  $u \in V$  do
     $d_s[u] \leftarrow \infty$ ;  $\pi_s[u] \leftarrow \text{null}$ 
 $d_s[s] \leftarrow 0$ ;
for  $i \leftarrow 1$  to  $|V|$  do
     $f \leftarrow \text{false}$ 
    foreach  $(u, v) \in E$  do
         $f \leftarrow f \vee \text{Relax}(u, v)$ 
    if  $f = \text{false}$  then return true
return false;
```