Introduction to Algorithms

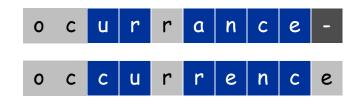
Dynamic Programming

Sequence Alignment

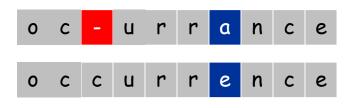
Word Alignment

How similar are two strings?

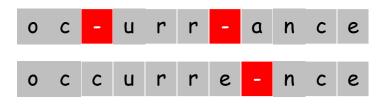
ocurrance
occurrence



5 mismatches, 1 gap



1 mismatch, 1 gap



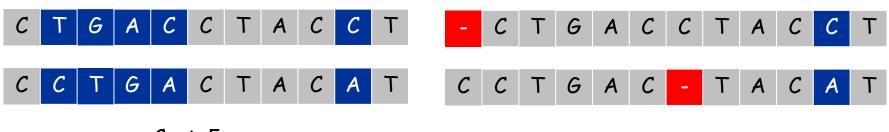
0 mismatches, 3 gaps

Edit Distance

Edit distance. [Levenshtein 1966, Needleman-Wunsch 1970] Cost = # of gaps + #mismatches.

Applications.

- Basis for Unix diff and Word correct in editors.
- Speech recognition.
- Computational biology.



Cost: 5

Sequence Alignment

Given two strings x_1 , ..., x_m and y_1 , ..., y_n find an alignment with minimum number of mismatch and gaps.

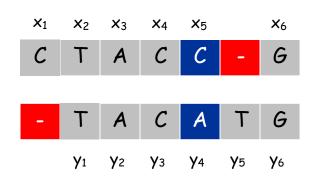
An alignment is a set of ordered pairs $(x_{i_1}, y_{j_1}), (x_{i_2}, y_{j_2}), \cdots$ such that $i_1 < i_2 < \cdots$ and $j_1 < i_2 < \cdots$

Example: CTACCG VS. TACATG.

Sol: We aligned

 x_2-y_1 , x_3-y_2 , x_4-y_3 , x_5-y_4 , x_6-y_6 .

So, the cost is 3.



DP for Sequence Alignment

Let OPT(i, j) be min cost of aligning x_i , ..., x_i and y_i , ..., y_j

Case 1: OPT matches x_i , y_j

• Then, pay mis-match cost if $x_i \neq y_j$ + min cost of aligning x_1 , ..., x_{i-1} and y_i , ..., y_{i-1} i.e., OPT(i-1, j-1)

Case 2: OPT leaves x_i unmatched

• Then, pay gap cost for $x_i + OPT(i-1, j)$

Case 3: OPT leaves y_j unmatched

• Then, pay gap cost for $y_i + OPT(i, j-1)$

Bottom-up DP

初始化与空字

符串的匹配

NEEDLEMAN-WUNSCH(T, S)

```
1: for i = 0 to m do
2: OPT[i, 0] = -3 * i;
```

3: end for

4: for
$$j = 0$$
 to n do

5:
$$OPT[0, j] = -3 * j;$$

6: end for

```
7: for j = 1 to n do
```

8: **for**
$$i=1$$
 to m **do**

9:
$$OPT[i, j] = \max\{OPT[i-1, j-1] + s(T_i, S_j), OPT[i-1, j] - 3, OPT[i, j-1] - 3\};$$

10: end for

11: end for

12: **return** OPT[m, n];

Analysis: $\Theta(mn)$ time and space.

English words or sentences: m, n \leq 10,...,20.

Computational biology: m = n = 100,000. 10 billions ops OK, but 40GB array?

	S:	1 1	0	C	U	R	R	A	N	C	E
occ oc_	T: ''	0	-3	-6	-9	-12	-15	-18	-21	-24	-27
	0	-3	1	-2	-5	-8	-11	-14	-17	-20	-23
	C	-6	-2	2	-1	-4	-7	-10	-13	-16	-19
	C	-9	-5	=1	1	-2	-5	-8	-11	-12	-15
	U	12	-8	-4	0	0	-3	-6	-9	-12	-13
	R	-15	-11	-7	-3	1	1	-2	-5	-8	-11
	R	-18	-14	-10	-6	-2	2	0	-3	-6	-9
	E	-21	-17	-13	-9	-5	-1	1	-1	-4	-5
	N	-24	-20	-16	-12	-8	-4	-2	2	-1	-4
	C	-27	-23	-19	-15	-11	-7	-5	-1	3	0
	E	-30	-26	-22	-18	-14	-10	-8	-4	0	4