

## 16 Sequence Alignment

- Sequence alignment
- Next: shortest paths in graphs with negative weights

### 16.1 Sequence Alignment

A way of comparing strings: line them up, possibly with gaps, to minimize the mismatches.

#### Example 16.1

Suppose we have strings *GACGTTA* and *GAACGCTA*.

We could align them as follows:

*GA \_CGTTA*  
*GAACGCTA*

But this is not a perfect alignment, since there are mismatches and gaps.

To quantify the quality of an alignment, we can define a **gap penalty**  $\delta$  and a **mismatch penalty**  $\alpha_{xy}$  for symbols  $x$  and  $y$ , with the total cost being the sum of the penalties.

Our goal is to find the alignment of lowest cost.

#### Example 16.2

*GA \_CG \_TTA*  
*GAACGCT \_A*

Here, the cost is  $3\delta$ .

#### Definition 16.3

Given strings  $x \in \Sigma^n$  and  $y \in \Sigma^m$  ( $\Sigma$  being an alphabet), a **matching** of sets  $I$  and  $J$  is a set of ordered pairs  $(i, j)$  with  $i \in I$ ,  $j \in J$  such that each  $i \in I$  and  $j \in J$  appears at most once (some can be unpaired).

#### Definition 16.4

A matching  $M$  of  $\{1, \dots, n\}$  and  $\{1, \dots, m\}$  is an **alignment** if there are no "crossings", i.e., if  $(i, j) \in M$  and  $(i', j') \in M$  with  $i < i'$ , then  $j < j'$ .

#### Example 16.5

$M = \{(1, 3), (2, 2)\}$  is a matching but not an alignment.

#### 16.1.1 Dichotomy for defining subproblems

Are the last two symbols matched? i.e., is  $(n, m) \in M$ ?

If not, then we know something more: either the last symbol of  $x$  or the last symbol of  $y$  must be unpaired. This is because if both  $x_n$  and  $y_m$  are matched, but not to each other, then there must be a crossing.

Subproblem: sequence alignment of  $x_1x_2 \dots x_i$  with  $y_1y_2 \dots y_j$ .

Let  $\text{opt}(i, j)$  be the minimum cost of an alignment between these substrings.

Is  $x_i$  aligned with  $y_j$ ?

- If yes, then  $\text{opt}(i, j) = \alpha_{x_i y_j} + \text{opt}(i-1)(j-1)$ .
- If no, then is  $x_i$  unmatched?

- If yes, then  $\text{opt}(i, j) = \delta + \text{opt}(i - 1, j)$
- If no, then  $y_j$  must be unmatched, so  $\text{opt}(i, j) = \delta + \text{opt}(i, j - 1)$ .

In general, we have

$$\text{opt}(i, j) = \min\{\alpha_{x_i y_j} + \text{opt}(i - 1, j - 1), \delta + \text{opt}(i - 1, j), \delta + \text{opt}(i, j - 1)\}$$

We define  $\text{opt}(i, 0) = i \cdot \delta$ , and  $\text{opt}(0, j) = j \cdot \delta$ .

Alignment (x, y):

```
let A be an (n+1)x(m+1) array indexed by
  i in {0, 1, ..., n} and j in {0, 1, ..., m}
let A[i, 0] = i * delta for all i in {0, 1, ..., n}
let A[0, j] = j * delta for all j in {0, 1, ..., m}
for i=1, ..., n
  for j=1, ..., m
    let A[i, j] = min{alpha_xiyj + A[i-1, j-1],
      delta + A[i-1, j], delta + A[i, j-1]}
  endfor
endfor
return A[n, m]
```