13 Sequence Alignment / Edit Distance [KT 6.6]

Definition. δ : gap penalty. $\alpha_{p,q}$: cost of matching $p,q \in \Sigma$ with $\alpha(p,p) = 0$ for all $p \in \Sigma$. We want to find an alignment between the two sequences $X \in \Sigma^m$ and $Y \in \Sigma^n$, such that the cost, which is measured by the *sum* of gap penalty and matching cost, is minimized.

Key observation. In an optimal alignment M, at least one of the following is true (m = |X|, n = |Y|). Proof by contradiction – otherwise it leads to crossing edges.

- 1. $(m, n) \in M$
- 2. X[m] is not matched
- 3. Y[n] is not matched

The recursion.

$$OPT(i, j) = \min\{\alpha_{x_i y_i} + OPT(i - 1, j - 1), \delta + OPT(i - 1, j), \delta + OPT(i, j - 1)\}.$$

Running time. Naive implementation needs O(mn) time and space. We only need to compute at most O(mn) of $OPT(\cdot, \cdot)$, and each can be computed in O(1) time (from previously already computed values).

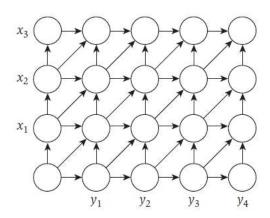


Figure 6.17 A graph-based picture of sequence alignment.

Equivalent to finding shortest path in a 2D grid. For example, to match 'mean' and 'name', suppose the gap penalty is $\delta = 2$, and $\alpha_{a,e} = \alpha_{m,n} = 1$, $\alpha_{a,m} = \alpha_{a,n} = \alpha_{e,m} = \alpha_{e,n} = 3$.

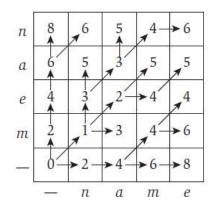


Figure 6.18 The opt values for the problem of aligning the words *mean* to *name*.

Better space (value only). Better implementation can improve the space to O(m+n). Easy to do from the shortest path representation: We compute the DP matrix row-by-row bottom-up, and at any time step we only keep the two most recent rows.

Why O(mn) space is *not* good? Since for example for DNA sequences, m and n can be very large. Say if they have 1,000,000 symbols, then the space would be 1 TB. On the other hand, if we have a $2.0 \mathrm{GHz}$ CPU, we can run an O(mn) time algorithm in about 500 seconds, which is a fairly reasonable time.

Find the actual alignment in O(m+n) space. The above O(m+n) space implementation cannot be used to find the optimal solution.

A better algo uses divide-and-conquer, and recursively finds the middle point in the shortest path, via a *bi-direction* dynamic programming.

Let f(i,j) denote the length of the shortest path from (0,0) to (i,j), and g(i,j) denote the length of the shortest path from (m,n) (back) to (i,j) (to better see this, one can rotate the DP matrix by 180 degree):

$$f(i,j) = \min\{\alpha_{x_i y_i} + f(i-1,j-1), \delta + f(i-1,j), \delta + f(i,j-1)\}.$$

$$g(i,j) = \min\{\alpha_{x_{i+1}y_{j+1}} + g(i+1,j+1), \delta + g(i+1,j), \delta + g(i,j+1)\}.$$

We have the following observation: The length of the shortest bottom-left corner to top-right corner path in the DP matrix that pass through (i,j) is f(i,j)+g(i,j). More generally, let $k \in \{0,\ldots,n\}$, and let $q \in \{0,\ldots,m\}$ be an index that minimize f(q,k)+g(q,k). Then there is a corner-to-corner path of minimum length that passes through the node (q,k). We then try out the q values one by one, and then find the one with the minimum cost. Finally, we recuse on two subproblems: to find the shortest path from (0,0) to (q,k), and that from (k,q) to (n,m). See the Figure 6.19 for an illustration.

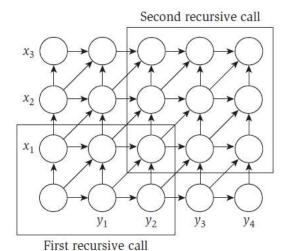


Figure 6.19 The first level of recurrence for the space-efficient Divide-and-Conquer-Alignment. The two boxed regions indicate the input to the two recursive cells.

The pseudocode of the algorithm is presented below.

Divide-And-Conquer-Alignment $(x_1, \ldots, x_n; y_1, \ldots, y_m)$

IF $(n \le 2)$ or $(m \le 2)$ THEN compute the best alignment using the old algorithm and stop; Let k = n/2;

Compute f[k,j], g[k,j] for all $j \in \{0,1,\ldots,m\}$ in time O(nm) and space O(n+m). Find a q such that f[k,q]+g[k,q] is minimized;

Insert the alignment path point (k, q) in the global path array;

Divide-And-Conquer-Alignment $(x_1, \ldots, x_k; y_1, \ldots, y_q);$

Divide-And-Conquer-Alignment $(x_{k+1}, \ldots, x_n; y_{q+1}, \ldots, y_m);$

The running time analysis of this algorithm is to solve the following recursion.

$$T(n,m) \le cmn + T(n/2,q) + T(n/2,m-q); T(2,m) \le cm; T(n,2) \le cn.$$

Solve the recursion by guessing $T(m, n) \leq \alpha mn$, and solve for $\alpha = 2c$.