

### **Dynamic Programming**

Some of the slides are courtesy of Charles Leiserson with small changes by Carola Wenk

#### Example: All-Pairs Shortest Paths Floyd-Warshall alg (Spring 2021)

- Given a graph G(V,E) with weights (positive and negative) assign to each edges. Assume V={v<sub>1</sub>...v<sub>n</sub>}.
- Compute a matrix D such that D[i,j] contains the length of the shortest path v<sub>i</sub> → v<sub>j</sub>
- Also compute a matrix ∏[1..n,1..n] such that ∏[i, j] is the vertex that proceed v<sub>i</sub> along the shortest path v<sub>i</sub> → v<sub>j</sub>
- Warshall-Floyd Algorithm computes these tables in O(n <sup>3</sup>)
- Can you think about alternative approaches when the weights of all edges is positive ?

In the figure to the right,  $k = \prod[i, j]$ .

Compare to  $\Pi[v_i]$  in Dijkstra or Bellman-Ford



#### **Dynamic Programming: Example 1: Longest Common Subsequance**

We look at sequences of characters (strings)

e.g. x = "ABCA"

**Def**: A **subsequence** of *x* is an sequence obtained from *x* by possibly deleting some of its characters (but without changing their order

Examples:<br/>"ABC","ACA","ABCA""ABC","ACA","ABCA"

**Def** A **prefix** of *x*, denoted x[1..m], is the sequence of the first *m* characters of *x* 

**Examples:**   $x[1..4] = "ABCA" \quad x[1..3] = "ABC" \quad x[1..2] = "AB"$  $x[1..1] = "A" \quad x[1..0] = ""$ 

#### Longest Common Subsequence (LCS)

• Given two sequences *x*[1 . . *m*] and *y*[1 . . *n*], find a longest subsequence common to them both.



Different phrasing: Find a set of a maximum number of segments, such that

•Each segment connects a character of x to an identical character of y,

- •Each character is used at most once
- •Segments do not intersect.

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# Cs445 salute



# **Brute-force LCS algorithm**

Checking every subsequence of x whether it is also a subsequence of y.

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### Analysis

- Checking =  $\Theta(m+n)$  time per subsequence.
- $2^m$  subsequences of x

Worst-case running time =  $\Theta$  ((*m*+*n*)2<sup>*m*</sup>) = exponential time.

# Towards a better algorithm

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# Towards a better algorithm

### Simplification:

- 1. Look at the *length* of a longest-common subsequence.
- 2. Extend the algorithm to find the LCS itself.

**Notation:** Denote the length of a sequence s by |s|.

**Strategy:** Consider *prefixes* of *x* and *y*.

• Define c[i, j] = |LCS(x[1 ... i], y[1 ... j])|.

• Then, c[m, n] = |LCS(x, y)|.

# **Recursive formulation**

Observation:

It is impossible that

x[m] is matched to an element in y[1..n-1] and simultaneously

*y*[*n*] is matched to an element in *x*[1..*m*-1] (since it must create a pair of crossing segments).

**Conclusion** – either x[m] is matched to y[n], or one at least of them is unmatched in **OPT**. {**OPT** – the optimal solution}



# **Recursive formaula**

Lets just consider the last character of of x and of y **Case (I):** x[m] = y[n]. Claim: c[m, n] = c[m-1, n-1]+1. *Proof.* 



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We claim that there is a max matching that matches x[m] to y[n].

Indeed, if x[m] is matched to y[k] (for k < m) then y[n] is unmatched (otherwise we have two crossing segments). Hence we can obtain another matching of the same cardinality by matching x[m] to y[n].

This implies that we can find an optimal matching of LCS(x[1..m-1] to y[1..n-1], and add the segment (x[m],y[n]).So c[m,n]=c[m-1,n-1]+1

## **Recursive formulation-cont**

**Case (II):**  $x[m] \neq y[n]$  Claim:  $c[m,n]=\max\{c[m,n-1], c[m-1,n]\}$ 

Recall - in LCS(x[1 ...m], y[1 ...n]) it cannot be that **both** x[m] and y[n] are both matched.



If x[m] is unmatched in OPT then LCS(x[1 ...m], y[1 ...n]) = LCS(x[1 ...m-1], y[1 ...n])If y[j] is unmatched in OPT then LCS(x[1 ...m], y[1 ...n]) = LCS(x[1 ...m], y[1 ...n-1])

So  $c[m,n] = \max\{c[m-1, n], c[m, n-1]\}$ 

## c[i,j] For general *i*,*j*

Since we only care for OPT matching the prefixes, then Case (I): x[i] = y[j]. Claim: if x[i] = y[j] then c[i, j] = c[i-1, j-1] + 1.

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**Optimal substructure** An optimal solution to a problem (instance) contains optimal solutions to subproblems.

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If z = LCS(x, y), then any prefix of z is an LCS of a prefix of x and a prefix of y.

# **Recursive algorithm for LCS**

```
LCS(x, y, i, j)

if ( i==0 or j=0) return 0

if x[i] = y[ j]

then return LCS(x, y, i-1, j-1) + 1

else return max{LCS(x, y, i-1, j), LCS(x, y, i, j-1)}
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To call the function LCS(x, y, m,n)

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Worst-case:  $x[i] \neq y[j]$ , for all *i*,*j* in which case the algorithm evaluates two subproblems, each with only one parameter decremented.





Height =  $m + n \Rightarrow$  work potentially  $2^{m+n}$  exponential.



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**Overlapping subproblems** A recursive solution contains a "small" number of distinct subproblems repeated many times.

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The number of distinct LCS subproblems for two strings of lengths m and n is only mn.

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   for j=0 to n c[0, j] = 0
   for i=1 to m
     for j=1 to n
        if (x[i] = y[j])
           then c[i, j] \leftarrow c[i-1, j-1] + 1
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```

Time =  $\Theta(mn)$  = constant work per table entry. Space =  $\Theta(mn)$ .

## LCS: Dynamic-programming algorithm



**IDEA:** Compute the table bottom-up. Fill *z* backward.

Observation:  $c[i;j] \ge c[i-1;j]$  and  $c[i;j] \ge c[i;j-1]$ **Proof Sketch:** We use a longer prefix, so there are more chars to be match.

#### LCS Reconstruction:

Set i=m; j=n; k=c[i;j]
While(k>0){
 if (c[i;j]>c[i-1;j] and c[i;j]>c[i;j-1]) {
 z[k] = x[i] ;
 i--; j--; k--;
 }else // c[i;j]=c[i-1;j] or c[i;j]=c[i-1;j]
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### Reconstructing *z=LCS(X,Y*)

Another idea – While filling c[], add arrows to each cell c[i,j] specifying which neighboring cell c[i,j] it got its value.

- *c[i,j].flag* = "\" if *c[i,,j]=c[i-1;j-1]+1*
- *c[i,j].flag* = "↑ " if *c[i,j]=c[i-1;j] c[i,j].flag* = "←" if *c[i,j]=c[i-1;j]*



# **Example 2: Edit distance**

Given strings X, Y, the edit distance ed(X, Y) between X and Y is defined as the minimum number of operations that we need to perform on X, in order to obtain Y.

**Definition**: An Operations (in this context) Insertion/Deletion/ Replacement of a **single** character.

#### Examples:

ed("aaba", "aaba")	= 0
ed("aaa", "aaba")	= 1
<i>ed("aaaa", "abaa")</i>	= 1
ed("baaa", "")	=4
<i>ed("baaa", "aaab")</i>	=2

Note that the term "distance" is a bit misleading: We need both the **value** (how many operations) as well as knowing **which** operations.

#### Example 3': "Priced" Edit distance *ed(X,Y)*

Assume also given

InsCost, - the cost of a single insertion into x.
DelCost - the cost of a single deletion from x, and
RepCost - the cost of replacing one character of x
by a different character.

**Definition:** Given strings X, Y, the **edit distance** ed(X, Y) between X and Y is the cheapest sequence of operations, starting on X and ending at Y.

**Problem:** Compute ed(X, Y), (both the value and the optimal sequence of operations.)

Definition: c[i,j] = Cost(ed(X[1..i], Y[1..j])).

Will first compute Cost( c[m,n]). Then will recover the sequence.

## Thm:

Let c[i,j] = ed(x[1..i], y[1..j]). Assume c[i-1,j-1], c[i-1,j-1], c[i-1,j] are already computed.

If X[i]=Y[j] then c[i,j] = c[i-1,j-1]Else //  $X[i] \neq Y[j]$   $c[i,j] = \min\{$   $c[i-1,j-1]+RepCost, //convert X[1..i-1] \rightarrow Y[1..j-1], \text{ and replace } y[j]$ by x[i]

 $\begin{array}{l} c[i-1, j] + DelCost, \ //delete \ X[i] \ and \ convert \ X[1..i-1] \rightarrow Y[1..j] \\ c[i,j-1] + InsCost \ //convert \ X[1..i,] \rightarrow Y[1..j-1], \ and \ insert \ Y[i] \\ \end{array}$ 

**Algorithm Memoization:** After computing a solution to a subproblem, store it in a table. Subsequent calls check the table to avoid redoing work.

## Algorithm

*Memoization:* After computing a solution to a subproblem, store it in a table. Subsequent calls check the table to avoid redoing work.

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ed(X, Y)
     for i=0 to m c[i, 0] = i DelCost
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           if (X[i] == Y[i])
               then c[i, j] \leftarrow c[i-1, j-1]
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```

Time =  $\Theta(m n)$  = constant work per table entry. Space =  $\Theta(m n)$ . Homework: Compute the sequence of operations. Compute which characters in *x* matches which chars in *y*.

## **Polygonal Path - definition**

- We definite a polygonal path  $P = \{p_1 \dots p_n\}$  where
- •Each vertex  $p_i$  is a point in the plane,
- •Vertex  $p_1$  is the first vertex,  $p_n$  is the last,
- •Vertex  $p_i$  is connected to the next vertex  $p_{i+1}$  by a straight segment.

$$\begin{array}{c|cccc} P & p_2 & p_4 \\ p_1 & p_3 & p_n \end{array}$$



- Should not be effected by how curves are sampled
- Should reflect the "order" of the points along the curves.



#### Problem: Computing the Frechet Distance between polylines *Frechet(P, Q,r)*

Definition of *Frechet(P,Q, r)* Assume a person walks on  $P = \{p_1 \dots p_n\}$ while a dog walks on  $Q = \{q_1 \dots q_n\}$ . *r* is the leash length (part of input). The **person** starts at  $p_1$  and ends at  $p_n$ The **dog** starts at  $q_1$  and ends at  $q_n$ 

 $q_2$ 

 $Q_{3}$ 

At each time stamp,

•either the **person** jumps to the next vertex

Or the dog jumps to the next vertexOr both jumps to the next vertex

• Every instance they stop, we measure whether the distance between person $\leftrightarrow$  dog (the length of the leash)  $\leq r$ .

 $p_n$ 

- Frechet(P,Q,r)=YES if the answer is positive for all time stamps.
- (if not, a longer leash is need. If yes, maybe a shorter one is sufficient.
- So we could use binary search.

#### **Problem:** Computing the Frechet Distance between polylines Frechet(**P**, **Q**, **r**) $q_2$ $p_n$ $Q_{3}$ Definition of *Frechet(P,Q, r)* Every instance they stop, we Assume a person walks on $P = \{p_1 \dots p_n\}$ • measure whether the distance while a dog walks on $Q = \{q_1, q_n\}$ . between person $\leftrightarrow$ dog (the length of the **leash**) $\leq r$ . *r* is the leash length (part of input). The **person** starts at $p_1$ and ends at $p_n$ Frechet(P,Q,r)=YES if the ulletThe dog starts at $q_1$ and ends at $q_n$ answer is positive for all time stamps.

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•Or the dog jumps to the next vertex

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 $Q_{z}$ 

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### **Computing Frechet(P,Q,r)**

Frechet(P,Q,r) // c[1..n, 1..n] - boolean array // *c[i,j]*=Frechet(P[1..*i*],Q[1..*j*], *r*)

Init:  $c[1,1] = (||p_1 - q_1|| \le r) (YES/NO)$ For i=2 to  $n c[i,1] = (||p_i - q_1|| \le r) AND c[i-1,1] (YES/NO)$ For j=2 to  $n c[1,j] = (||p_1 - q_i|| \le r) AND c[1,j-1]$ 

### Computing Frechet (P,Q,r) (cont.)

// c[1..n, 1..n] – boolean array

Init- previous slide

For i=2 to nFor j=2 to n  $c[i,j] = (||p_i - q_j|| \le r)$  AND  $\{ c[i-1,j-1], // both jumps$ OR  $c[i-1,j], // person jumped from <math>p_{i-1}$  to  $p_i$ , dog stays at  $q_j$ OR c[i,j-1]. // person stayed at  $p_i$ , dog jumped from  $q_{j-1}$  to  $q_j$ .  $\}$ 

**Return** *c[n.n]* 

Note – this is only the cost (that is the distance itself. We still need to find what is the series of steps that yield this cost

### Comments

- This is actually the **Discrete** Frechet Distance (only distances between vertices counts). We do not discuss the **continuous** version.
- This is only the Decision problem we actually want the shortest leash. We could use a binary search to approximate it. Exact algorithm outside the scope of this course
- If person/dog could move backward, the problem is called the **weak** Frechet.



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The input is the locations of their vertices (e.g. GIS coordinates)

How similar are P to Q?

Need to come up with a number dtw(P,Q)? So if dtw(P,Q) < dtw(P,Q'), then **P** is more similar to **Q** 



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# **Dynamic Time Warping** dtw(P,Q) $P \stackrel{p_2}{q_2} \qquad p_4$ $p_1 \stackrel{p}{q_2} \qquad p_3$ $q_3$

Definition of dtw(P,Q)Assume a person walks on  $P = \{p_1 \dots p_n\}$ while a dog walks on  $Q = \{q_1 \dots q_m\}$ .

They **person** starts at  $p_1$  and ends at  $p_n$ They **dog** starts at  $q_1$  and ends at  $q_n$ 

At each time stamp,

•either the **person** jumps to the next vertex

Or the dog jumps to the next vertexOr both jumps to the next vertex

- Every instance they stop,
   we measure the distance
   (the length of the leash)
   person↔dog.
- We sum the lengths of all leashes.
- dtw(P,Q) is the smallest sum (over all possible sequences)

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 $Q_2$ 

 $q_3$ 

**Dynamic Time Warping** *dtw(P,Q)* 

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#### Motivation:



Definition of dtw(P,Q)Assume a person walks on  $P = \{p_1 \dots p_n\}$ while a dog walks on  $Q = \{q_1 \dots q_m\}$ .

Distance between trajectoris enables finding nearest neighbor, and clustering

But two very similar trajectories might have vertices in very different places DTW is used in

- Signal processing (speech reco)
- Signature verification
- Analysis of vehicles trajectories for roads networks
- Improving locationsbased services
- Animals migrations patters
- Stocks analysis

## **Thm 1:**

Let c[i,j] = dtw(P[1..i], Q[1..j]).

Let  $|| p_i - q_j ||$  be the between the points  $p_i$  and  $q_j$ That is, the length of the leash.

For every i > 1, j > 1 $c[1,1] = || p_1 - q_1 ||$ 

 $c[1,j] = c[1,j-1] + ||p_1 - q_j||$ 

 $c[i,1] = c[i-1,1] + ||p_i - q_1||$ 

## **Thm 2**:

Assume at some time, the person is at  $p_i$  while dog at  $q_{j}$ . Assume i > 1 and j > 1.

What (might have) happened one step ago ?

Three possibilities

Both person and the dog jumped (from  $p_{i-1}$  and from  $q_j$ ) OR Person jumped from  $p_{i-1}$  to  $p_i$ , dog stays at  $q_j$  OR Person stayed at  $p_i$ , dog jumped from  $q_{i-1}$  to  $q_{i-1}$ 

## Thm 2 cont:

Let c[i,j] = dtw(P[1..i], Q[1..j]).

If  $i \ge 1$  and  $j \ge 1$  then

 $c[i,j] = || p_i - q_j || + \min\{c[i-1,j-1], // both jumps c[i-1,j], // both jumps c[i-1,j], // person jumped from p_{i-1} to p_i, dog stays at q_j c[i,j-1]. // person stayed at p_i, dog jumped from q_{j-1} to q_{j-1} \}$ 

Since we are not sure that when the person is at  $p_i$  the dog is at  $q_j$  we will compute all such pairs i, j – one of them must happened

#### Algorithm for computing dtw(P,Q) Init according to Thm 1.

```
Fot i=2 to n

For j=2 to n

c[i,j] = || p_i - q_j || + \min\{c[i-1,j-1], // both jumps c[i-1,j], // person jumped from <math>p_{i-1} to p_i, dog stays at q_j

c[i,j-1] // person stayed at <math>p_i, dog jumped from q_{j-1} to q_j.

}

Peturn c[n, n]
```

#### **Return** *c*[*n*.*n*]

Note – this is only the cost (that is the distance itself. We still need to find what is the series of steps that yield this cost

#### **Dynamic-programming hallmark #1**

(we saw this slide already)

**Optimal substructure** An optimal solution to a problem (instance) contains optimal solutions to subproblems.

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**Optimal substructure** An optimal solution to a problem (instance) contains optimal solutions to subproblems.

If z = LCS(x, y), then any prefix of z is an LCS of a prefix of x and a prefix of y.

### **Dynamic-programming hallmark** #2

**Overlapping subproblems** A recursive solution contains a "small" number of distinct subproblems repeated many times.

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**Overlapping subproblems** A recursive solution contains a "small" number of distinct subproblems repeated many times.

The number of distinct LCS subproblems for two strings of lengths m and n is only mn.

#### Another application of DP: **Clustering** (source: Kleinberg & Tardos)



$$\frac{2g_i}{n}$$



• Given points  $P = (p_1, p_2, \dots, p_n)$  sorted from left to right, and a panelty R, find optimal k, and partition of P into k **runs**  $(p_1, p_2 \dots p_{i_1})(p_{i_1+1}, p_{i_1+2} \dots p_{i_2}), (p_{i_2+1}, \dots, p_{i_3}) \dots (p_{i_{k-1}+1} \dots p_n)$ and lines  $\ell_1 \dots \ell_k$  (one per each run) So that the sum  $R + Err(\ell_1, \{p_1, p_2 \dots p_{i_1}\}) +$ 

$$R + Err(\ell_2, \{p_{i_1+2} \dots p_{i_2}\}) +$$

$$R + Err(\ell_k, \{p_{i_{k-1}+1} \dots p_n\})$$

is as small as possible

÷

Note that if R=0, we will probably use n/2 runs  $(p_1 p_2)$ ,  $(p_3, p_4)$ , ...  $(p_{n-1}, p_n)$ . If R is huge, we can afford only one penalty, so only one run  $(p_1...,p_n)$ . In the example, k=3,  $i_1=5$ ,  $i_2=8$ 



- Algorithm:
- Preprocessing:  $\forall j < i$ : compute the line  $\ell$  minimizing the error for the set  $\{p_j, p_{j+1} \dots p_i\}$ .

Let 
$$e[j, i] = Err(\ell, \{p_j, p_{j+1} \dots p_i\})$$

- Idea: Let c[i] = cost of the opt clustering problem for the set  $\{p_1 \dots p_i\}$ .
- Init: c[0] = 0.
- for i = 2 to n do {

  c[i] = min{R + c[j] + e[j + 1, i] | 0 ≤ j < i}</li>

  return c[n]
## Summarizing

- The algorithm takes  $O(n^3)$  and  $O(n^2)$  space
- (for preprocessing *d[j,i]* )
- Note we did not discuss how to reconstruct the solution itself. We only calculated its cost