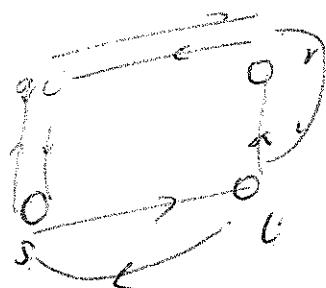


Elements of Dynamic Programming

- Optimal substructure
- Bottom up
- Subproblems share a parent node



- Longest path does not have independent subproblems.
- Overlapping subproblems, Reusing results in optimal min
- Longest common subsequence

- Biological & genomic applications

- Subsequence, $X = \langle x_1, x_2, \dots, x_m \rangle$
- Common subsequence $Z = \langle z_1, z_2, \dots, z_k \rangle$
- Common subsequence $\langle c_1, c_2, \dots, c_k \rangle$ c_i, c_{i+1}, \dots, c_k
- Form $x_{ij} = z_0$

- longest common subsequence

- Brute force approach

- Optimal Substructure

$$\text{Let } X_i = \langle x_1, x_2, \dots, x_i \rangle \quad x_0 = \epsilon$$

example

Thm :- Let $X = \langle x_1, \dots, x_m \rangle \quad Y = \langle y_1, \dots, y_n \rangle$
 $Z = \langle z_1, z_2, \dots, z_k \rangle$ be an LCS of X and Y

If $x_m = y_n$, then $z_k = x_m = y_n \in Z_{k-1} = \text{LCS}(X_{m-1}, Y_{n-1})$

If $x_m \neq y_n$ and $z_k \neq x_m \rightarrow Z = \text{LCS}(X_{m-1}, Y_n)$

If $x_m \neq y_n$ and $z_k \neq y_n \rightarrow Z = \text{LCS}(X, Y_{n-1})$

Let $c(i,j) = \text{length of LCS of sequences } X_i \text{ and } Y_j$

$$c(i,j) = \begin{cases} 0, & \text{if } i=0, j=0 \\ c(i-1, j-1), & \text{if } i, j > 0 \text{ and } x_i = y_j \\ \max(c(i-1, j), c(i, j-1)), & \text{if } i, j > 0 \text{ &} \\ & x_i \neq y_j \end{cases}$$

y_j	B	D	C	A	B	A	
x_i	0	0	0	0	0	0	
	0	0	0	1	1	1	
A	0	0	0	1	1	1	
B	1	1	1	1	2	2	
C	2	2	2	2	2	2	
	0	1	1	2	2	2	
	1	1	2	2	2	2	
	2	2	2	2	2	2	
	3	3	3	3	3	3	
	4	4	4	4	4	4	
	5	5	5	5	5	5	

If $c(i-1, j)$

$\geq c(i, j-1)$

$b(i, j) = \uparrow$

else

$b(i, j) = \leftarrow$

y_j	B	D	C	A	B	A	
x_i	0	0	0	0	0	0	
	0	0	0	1	1	1	
A	0	0	0	1	1	1	
B	1	1	1	1	2	2	
C	2	2	2	2	2	2	
	0	1	1	2	2	2	
	1	1	2	2	2	2	
	2	2	2	2	2	2	
	3	3	3	3	3	3	
	4	4	4	4	4	4	
	5	5	5	5	5	5	