Homework 1

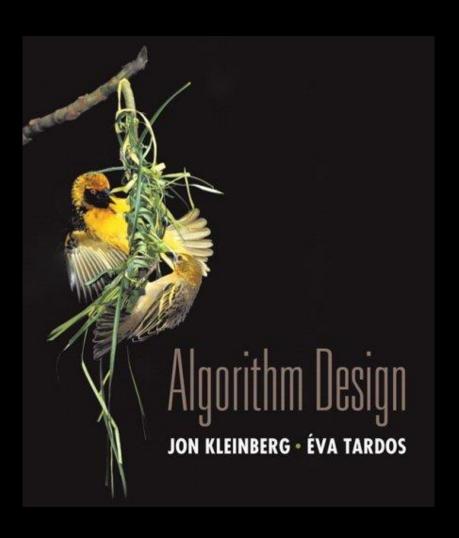
Homework 1

- Covers chapters 2, 4, 5, 6
- To be released today in Blackboard
- Due: 11:59pm, Mar. 23
- No late homework will be accepted!



Submission

- Register at <u>gradescope.com</u> using your university email
 - Make sure that your FULL NAME is your Chinese name
- Enroll in this course with code 968Y6E
- Save your homework solutions in a PDF or image file
- Upload your file to Gradescope and match your solution to each problem



Chapter 6

Dynamic Programming



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Algorithmic Paradigms

按照一个序列的顺序来选择

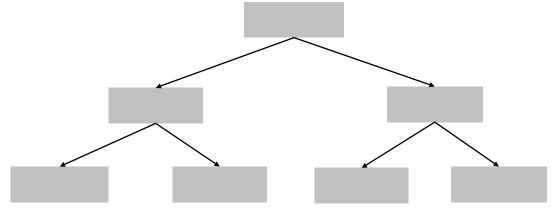
Greed. Build up a solution incrementally, myopically optimizing some local criterion.

Divide-and-conquer. Break up a problem into a few sub-problems, solve each sub-problem independently and recursively, and combine solution to sub-problems to form solution to original problem.

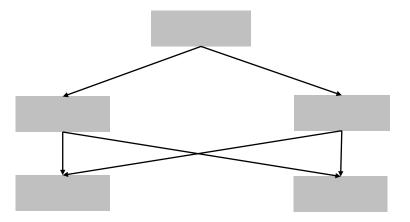
Dynamic programming. Break up a problem into a series of overlapping sub-problems, and build up solutions to larger and larger sub-problems.

Divide-and-conquer VS. Dynamic programming

Divide-and-conquer



Dynamic programming

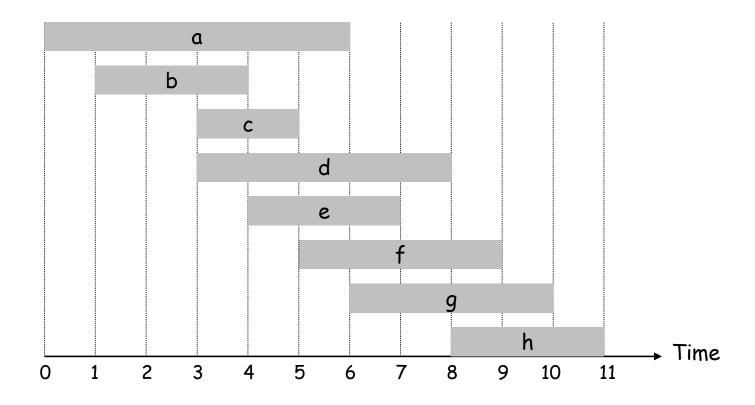


6.1 Weighted Interval Scheduling

Weighted Interval Scheduling

Weighted interval scheduling problem.

- \blacksquare Job j starts at s_j , finishes at f_j , and has weight or value v_j .
- Two jobs compatible if they don't overlap.
- Goal: find maximum weight subset of mutually compatible jobs.

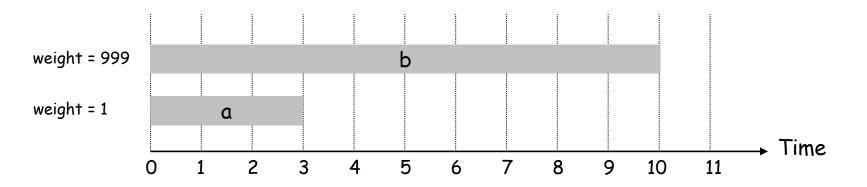


Unweighted Interval Scheduling Review

Recall. Greedy algorithm works if all weights are 1.

- Consider jobs in ascending order of finish time.
- Add job to subset if it is compatible with previously chosen jobs.

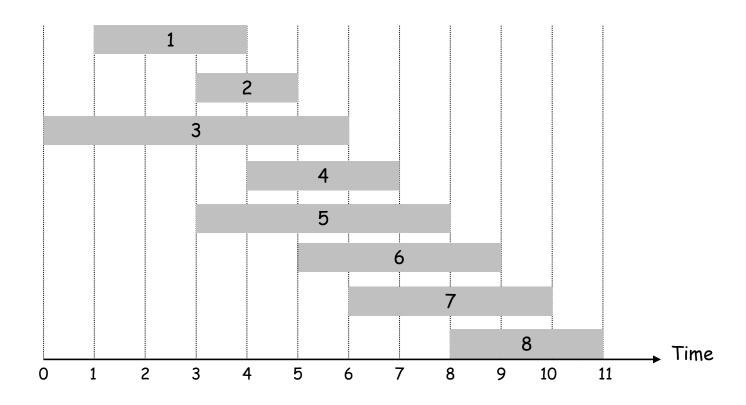
Observation. Greedy algorithm can fail spectacularly if arbitrary weights are allowed.



Weighted Interval Scheduling

Notation. Label jobs by finishing time: $f_1 \le f_2 \le ... \le f_n$. Def. p(j) = largest index i < j such that job i is compatible with j.

Ex:
$$p(8) = 5$$
, $p(7) = 3$, $p(2) = 0$.



Dynamic Programming: Binary Choice

Notation. OPT(j) = value of optimal solution to the problem consisting of job requests 1, 2, ..., j.

- Case 1: OPT selects job j.
 - can't use incompatible jobs { p(j) + 1, p(j) + 2, ..., j 1 }
 - must include optimal solution to problem consisting of remaining compatible jobs 1, 2, ..., p(j)

 optimal substructure
- Case 2: OPT does not select job j.
 - must include optimal solution to problem consisting of remaining compatible jobs 1, 2, ..., j-1

$$OPT(j) = \begin{cases} 0 & \text{if } j = 0\\ \max \{ v_j + OPT(p(j)), OPT(j-1) \} & \text{otherwise} \end{cases}$$

$$Case 1 \qquad Case 2$$

Weighted Interval Scheduling: Brute Force

Brute force algorithm.

```
Input: n, s_1,...,s_n, f_1,...,f_n, v_1,...,v_n

Sort jobs by finish times so that f_1 \leq f_2 \leq ... \leq f_n.

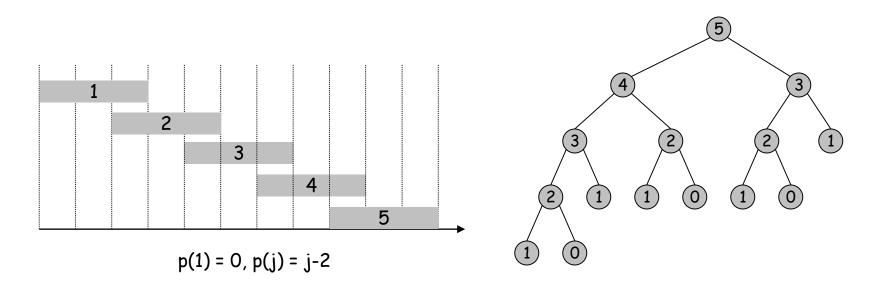
Compute p(1), p(2), ..., p(n)

Compute-Opt(j) {
   if (j = 0)
     return 0
   else
     return max(v_j + Compute-Opt(p(j)), Compute-Opt(j-1))
}
```

Weighted Interval Scheduling: Brute Force

Observation. Recursive algorithm fails spectacularly because of redundant sub-problems \Rightarrow exponential algorithms.

Ex. Number of recursive calls for family of "layered" instances grows like Fibonacci sequence.



Weighted Interval Scheduling: Memoization

Memoization. Store results of each sub-problem in a cache; lookup as needed.

```
Input: n, s_1, ..., s_n, f_1, ..., f_n, v_1, ..., v_n
Sort jobs by finish times so that f_1 \le f_2 \le \ldots \le f_n.
Compute p(1), p(2), ..., p(n)
for j = 1 to n
   M[j] = empty \leftarrow global array
M[0] = 0
M-Compute-Opt(j) {
   if (M[j] is empty)
       M[j] = max(w_j + M-Compute-Opt(p(j)), M-Compute-Opt(j-1))
   return M[j]
```

Weighted Interval Scheduling: Running Time

Claim. Memoized version of algorithm takes O(n log n) time.

- Sort by finish time: O(n log n).
- Computing $p(\cdot)$: O(n) after sorting by start time \leftarrow how?
- M-Compute-Opt(j): O(n)
 - Each entry M[j] is computed only once
 - The computation of M[j] invokes M-Compute-Opt twice

Remark. O(n) if jobs are pre-sorted by start and finish times.

Weighted Interval Scheduling: Finding a Solution

- Q. Dynamic programming algorithms computes optimal value. What if we want the solution itself?
- A. Do some post-processing.

```
Run M-Compute-Opt(n)
Run Find-Solution(n)

Find-Solution(j) {
   if (j = 0)
      output nothing
   else if (v<sub>j</sub> + M[p(j)] > M[j-1])
      print j
      Find-Solution(p(j))
   else
      Find-Solution(j-1)
}
```

• # of recursive calls \leq n \Rightarrow O(n).

Weighted Interval Scheduling: Bottom-Up

Bottom-up dynamic programming. Unwind recursion.

```
Input: n, s_1,...,s_n, f_1,...,f_n, v_1,...,v_n

Sort jobs by finish times so that f_1 \leq f_2 \leq ... \leq f_n.

Compute p(1), p(2), ..., p(n)

Iterative-Compute-Opt {

M[0] = 0

for j = 1 to n

M[j] = max(v_j + M[p(j)], M[j-1])
}
```

Top-down vs. bottom-up

- Top-down: May skip unnecessary sub-problems
- Bottom-up: Save the overhead in recursion

6.4 Knapsack Problem

Knapsack Problem

Knapsack problem.

- Given n objects and a "knapsack."
- Item i weighs $w_i > 0$ kilograms and has value $v_i > 0$.
- Knapsack has capacity of W kilograms.
- Goal: fill knapsack so as to maximize total value.



Knapsack Problem

Knapsack problem.

- Given n objects and a "knapsack."
- Item i weighs $w_i > 0$ kilograms and has value $v_i > 0$.
- Knapsack has capacity of W kilograms.
- Goal: fill knapsack so as to maximize total value.

Ex: { 3, 4 } has value 40.

W = 11

Item	Value	Weight
1	1	1
2	6	2
3	18	5
4	22	6
5	28	7

Greedy:

- repeatedly add item with maximum value v_i.
- lacksquare repeatedly add item with maximum weight w_i .
- repeatedly add item with maximum ratio v_i / w_i .

Greedy not optimal!

Dynamic Programming: False Start

Def. OPT(i) = max profit subset of items 1, ..., i.

- Case 1: OPT does not select item i.
 - OPT selects best of { 1, 2, ..., i-1 }
- Case 2: OPT selects item i.
 - How shall we enforce the weight limit? 需要提供重量/限制的信息

Conclusion. Shall specify the remaining weight capacity in OPT

Dynamic Programming: Adding a New Variable

Def. OPT(i, w) = max profit subset of items 1, ..., i with weight limit w.

- Case 1: OPT does not select item i.
 - OPT selects best of { 1, 2, ..., i-1 } using weight limit w
- Case 2: OPT selects item i.
 - new weight limit = w wi
 - OPT selects best of { 1, 2, ..., i-1 } using this new weight limit

$$OPT(i,w) = \begin{cases} 0 & \text{if } i = 0 \\ OPT(i-1,w) & \text{if } w_i > w \\ \max \left\{ OPT(i-1,w), \quad v_i + OPT(i-1,w-w_i) \right\} & \text{otherwise} \\ \text{最优子结构}, \ \texttt{包含前i-thmap} \\ \text{最优子结构}, \ \texttt{包含前i-thmap} \\ \end{cases}$$

一定要找出当前状态和之前状态的关系

Knapsack Problem: Bottom-Up

Knapsack. Fill up an n-by-W array.

```
Input: n, w_1, ..., w_N, v_1, ..., v_N
for w = 0 to W
  M[0, w] = 0
for i = 1 to n
   for w = 1 to W
      if (w_i > w)
          M[i, w] = M[i-1, w]
      else
          M[i, w] = \max \{M[i-1, w], v_i + M[i-1, w-w_i]\}
return M[n, W]
```

Knapsack Algorithm

		0	1	2	3	4	5	6	7	8	9	10	11
	ф												
	{ 1 }												
n + 1	{ 1, 2 }												
	{1,2,3}												
	{ 1, 2, 3, 4 }												

Item	Value	Weight
1	1	1
2	6	2
3	18	5
4	22	6
5	28	7

{1,2,3,4,5}

$$OPT(i, w) = \begin{cases} 0 & \text{if } i = 0 \\ OPT(i-1, w) & \text{if } w_i > w \\ \max \{ OPT(i-1, w), v_i + OPT(i-1, w-w_i) \} & \text{otherwise} \end{cases}$$

$$W = 11$$

Knapsack Algorithm

_____ W + 1 _____

		0	1	2	3	4	5	6	7	8	9	10	11
	ф	0	0	0	0	0	0	0	0	0	0	0	0
	{ 1 }												
n + 1	{ 1, 2 }												
	{1,2,3}												
	{1,2,3,4}												
	{1,2,3,4,5}												

Item	Value	Weight
1	1	1
2	6	2
3	18	5
4	22	6
5	28	7

$$OPT(i, w) = \begin{cases} 0 & \text{if } i = 0 \\ OPT(i-1, w) & \text{if } w_i > w \\ \max \{ OPT(i-1, w), v_i + OPT(i-1, w-w_i) \} & \text{otherwise} \end{cases}$$

$$W = 11$$

Knapsack Algorithm

		0	1	2	3	4	5	6	7	8	9	10	11	
	ф	0	0	0	0	0	0	0	0	0	0	0	0	i=0
	{ 1 }	0	1	1	1	1	1	1	1	1	1	1	1	i=1
n + 1	{ 1, 2 }	0_	1	6 _{Ca}	7 se 2	7	7	7	7	7	7	7	7	
	{ 1, 2, 3 }	0	1	6	7	7	18_	19	24	25 Cas	25	25	25	
	{1,2,3,4}	0	1	6	7	7	18	22	24	28	29	29	4 0 <i>C</i> a	co 1
	{1,2,3,4,5}	0	1	6	7	7	18	22	28	29	34	34	40	36 I

Item	Value	Weight
1	1	1
2	6	2
3	18	5
4	22	6
5	28	7

$$OPT(i, w) = \begin{cases} 0 & \text{if } i = 0 \\ OPT(i-1, w) & \text{if } w_i > w \\ \max \{ OPT(i-1, w), v_i + OPT(i-1, w-w_i) \} & \text{otherwise} \end{cases}$$

W = 11

Knapsack Algorithm: Top-down

		0	1	2	3	4	5	6	7	8	9	10	11	
	ф		0	0	0	0	0	0		0	0	0	0	
	{ 1 }			1	1	1	1	1			1		1	
n + 1	{ 1, 2 }	0				7	7	7					7	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \
	{ 1, 2, 3 }					7	18							为了知 道那些
	{ 1, 2, 3, 4 }					7								数值需 要计算
	{1,2,3,4,5}												40	

Item	Value	Weight
1	1	1
2	6	2
3	18	5
4	22	6
5	28	7

$$OPT(i, w) = \begin{cases} 0 & \text{if } i = 0 \\ OPT(i-1, w) & \text{if } w_i > w \\ \max \{ OPT(i-1, w), v_i + OPT(i-1, w-w_i) \} & \text{otherwise} \end{cases}$$

$$W = 11$$

Knapsack Problem: Running Time

Running time. $\Theta(n W)$.

- Not polynomial in input size!
- "Pseudo-polynomial."
- Decision version of Knapsack is NP-complete. [Chapter 8]

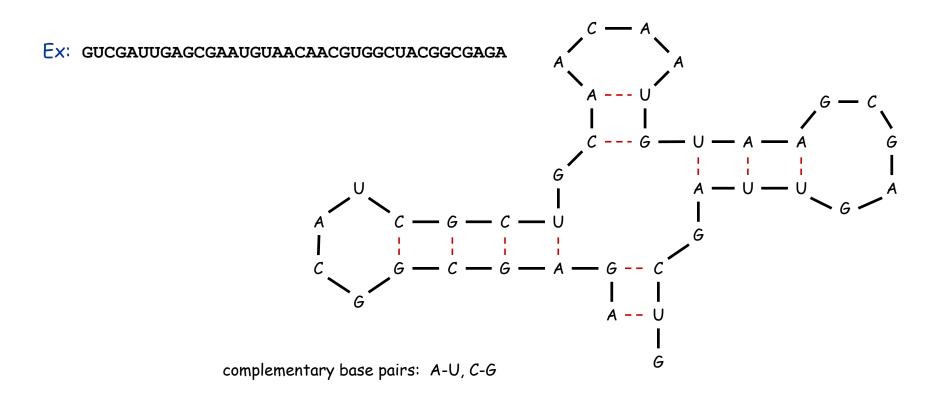
Knapsack approximation algorithm. There exists a polynomial algorithm that produces a feasible solution that has value within 0.01% of optimum. [Section 11.8]

6.5 RNA Secondary Structure

RNA Secondary Structure

RNA. String B = $b_1b_2...b_n$ over alphabet { A, C, G, U }.

Secondary structure. RNA is single-stranded so it tends to loop back and form base pairs with itself. This structure is essential for understanding behavior of molecule.



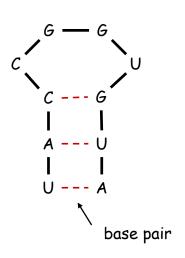
RNA Secondary Structure

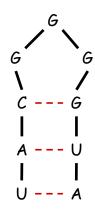
Secondary structure. A set of pairs $S = \{(b_i, b_j)\}$ that satisfy:

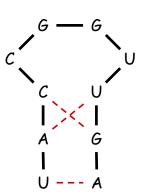
- [Watson-Crick.] S is a matching and each pair in S is a Watson-Crick complement: A-U, U-A, C-G, or G-C.
- [No sharp turns.] The ends of each pair are separated by at least 4 intervening bases. If $(b_i, b_j) \in S$, then i < j 4.
- [Non-crossing.] If (b_i, b_j) and (b_k, b_l) are two pairs in S, then we cannot have i < k < j < l.

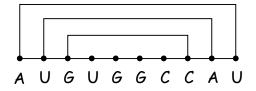
RNA Secondary Structure: Examples

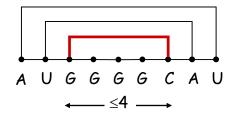
Examples.

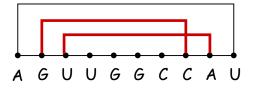












ok

sharp turn

crossing

RNA Secondary Structure

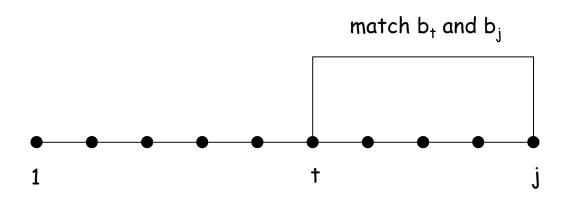
Free energy. Usual hypothesis is that an RNA molecule will form the secondary structure with the optimum total free energy.

approximate by number of base pairs

Goal. Given an RNA molecule $B = b_1b_2...b_n$, find a secondary structure S that maximizes the number of base pairs.

RNA Secondary Structure: Subproblems

First attempt. OPT(j) = maximum number of base pairs in a secondary structure of the substring $b_1b_2...b_j$.



Difficulty. Results in two sub-problems.

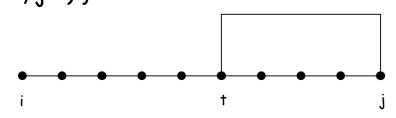
- Finding secondary structure in: $b_1b_2...b_{t-1}$. $\leftarrow OPT(t-1)$
- Finding secondary structure in: $b_{t+1}b_{t+2}...b_{i-1}$. \leftarrow need more sub-problems

Dynamic Programming Over Intervals

Notation. OPT(i, j) = maximum number of base pairs in a secondary structure of the substring $b_i b_{i+1} ... b_j$.

- If $i \ge j 4$.
 - OPT(i, j) = 0 by no-sharp turns condition.
- If i < j-4: take max of two cases</p>
 - Case 1. Base b_j is not involved in a pair. OPT(i, j-1)
 - Case 2. Base b_j pairs with b_t for some $i \le t < j 4$. Non-crossing constraint decouples resulting sub-problems $1 + \max_t \{ OPT(i, t-1) + OPT(t+1, j-1) \}$

take max over t such that $i \leq t < j\text{-}4$ and b_t and b_j are Watson-Crick complements

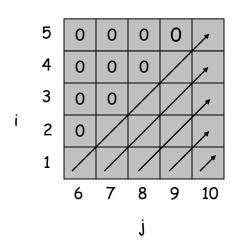


Bottom Up Dynamic Programming Over Intervals

- Q. What order to solve the sub-problems?
- A. Do shortest intervals first.

```
RNA(b<sub>1</sub>,...,b<sub>n</sub>) {
  for k = 5, 6, ..., n-1
    for i = 1, 2, ..., n-k
        j = i + k
        Compute M[i, j]

return M[1, n]
}
```



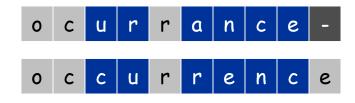
Running time. $O(n^3)$.

6.6 Sequence Alignment

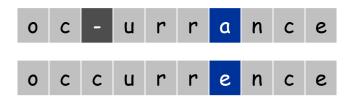
String Similarity

How similar are two strings?

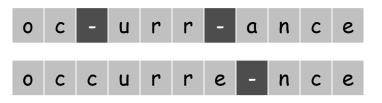
- ocurrance
- occurrence



6 mismatches, 1 gap



1 mismatch, 1 gap

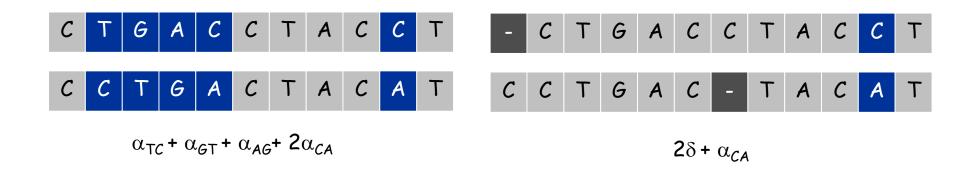


0 mismatches, 3 gaps

Edit Distance

Edit distance. [Levenshtein 1966, Needleman-Wunsch 1970]

- Gap penalty δ ; mismatch penalty α_{pq} .
- Cost = sum of gap and mismatch penalties.
- Edit distance = min cost



Applications.

- Basis for Unix diff.
- Speech recognition.
- Computational biology.

Sequence Alignment

Goal: Given two strings $X = x_1 x_2 ... x_m$ and $Y = y_1 y_2 ... y_n$ find alignment of minimum cost.

Def. An alignment M is a set of ordered pairs x_i - y_j such that each item occurs in at most one pair and no crossings.

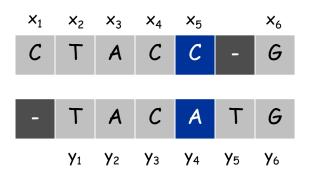
• The pair $x_i - y_j$ and $x_{i'} - y_{j'}$ cross if i < i', but j > j'.

Def. The cost of an alignment

$$cost(M) = \underbrace{\sum_{(x_i, y_j) \in M} \alpha_{x_i y_j}}_{\text{mismatch}} + \underbrace{\sum_{i: x_i \text{ unmatched } j: y_j \text{ unmatched}}_{\text{gap}} \delta$$

Ex: An alignment of CTACCG vs. TACATG.

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



Sequence Alignment: Problem Structure

Def. OPT(i, j) = min cost of aligning strings $x_1 x_2 ... x_i$ and $y_1 y_2 ... y_j$.

- Case 1: OPT matches x_i-y_j .
 - pay mismatch for x_i - y_j + min cost of aligning two strings $x_1 x_2 \dots x_{i-1}$ and $y_1 y_2 \dots y_{j-1}$
- Case 2a: OPT leaves x_i unmatched.
 - pay gap for x_i and min cost of aligning $x_1 x_2 \dots x_{i-1}$ and $y_1 y_2 \dots y_j$
- Case 2b: OPT leaves y_i unmatched.
 - pay gap for y_j and min cost of aligning $x_1 x_2 \dots x_i$ and $y_1 y_2 \dots y_{j-1}$

$$OPT(i, j) = \begin{cases} j\delta & \text{if } i = 0 \\ \alpha_{x_i y_j} + OPT(i-1, j-1) & \text{otherwise} \\ \delta + OPT(i, j-1) & \text{otherwise} \end{cases}$$

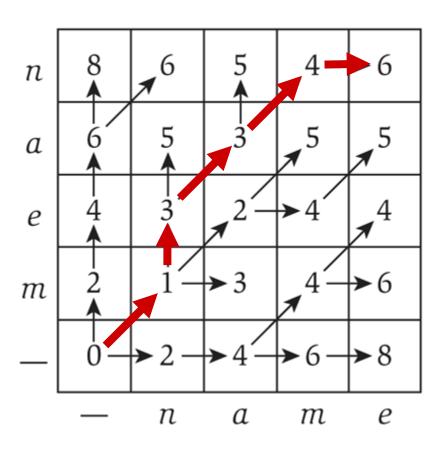
$$i\delta & \text{if } j = 0$$

Sequence Alignment: Algorithm

```
Sequence-Alignment(m, n, x_1x_2...x_m, y_1y_2...y_n, \delta, \alpha) {
   for i = 0 to m
      M[0, i] = i\delta
   for j = 0 to n
      M[j, 0] = j\delta
   for i = 1 to m
       for j = 1 to n
          M[i, j] = min(\alpha[x_i, y_j] + M[i-1, j-1],
                            \delta + M[i-1, j],
                            \delta + M[i, j-1]
   return M[m, n]
```

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

Sequence Alignment: Example



Sequence Alignment: Algorithm

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

English words or sentences:

■ m, n ≤ 30. ← OK

Computational biology:

- m = n = 100,000
- 10 billions ops is OK, but 10GB array is quite large

6.7 Sequence Alignment in Linear Space

Q. Can we avoid using quadratic space?

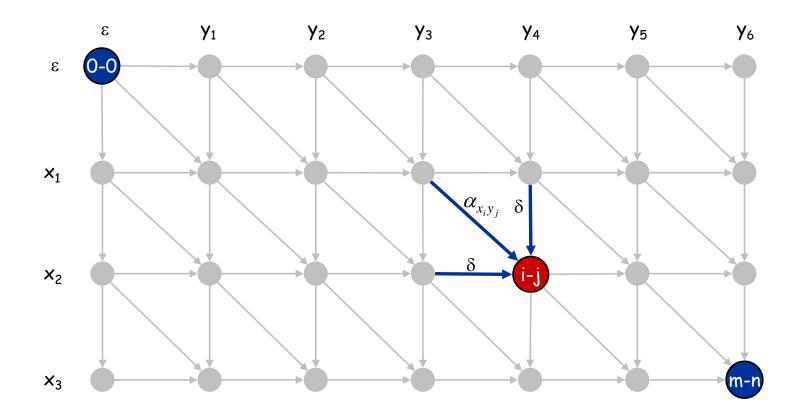
Easy. Optimal cost in O(m + n) space and O(mn) time.

- Compute OPT(i, •) from OPT(i-1, •).
- No longer a simple way to recover alignment itself.

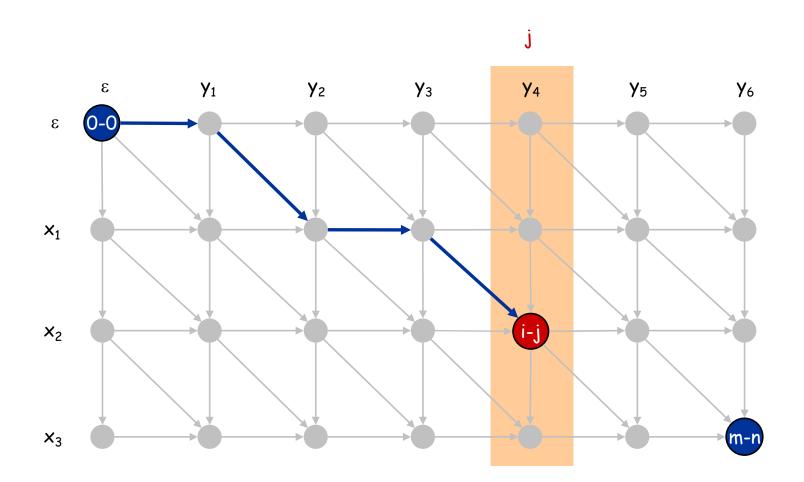
Theorem. [Hirschberg 1975] Optimal alignment in O(m + n) space and O(mn) time.

Clever combination of divide-and-conquer and dynamic programming.

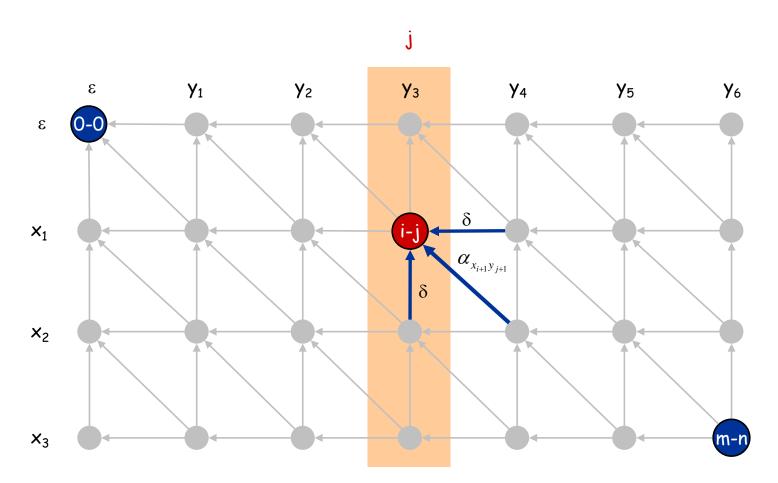
- Let f(i, j) be shortest path from (0,0) to (i, j).
- Observation: f(i, j) = OPT(i, j).



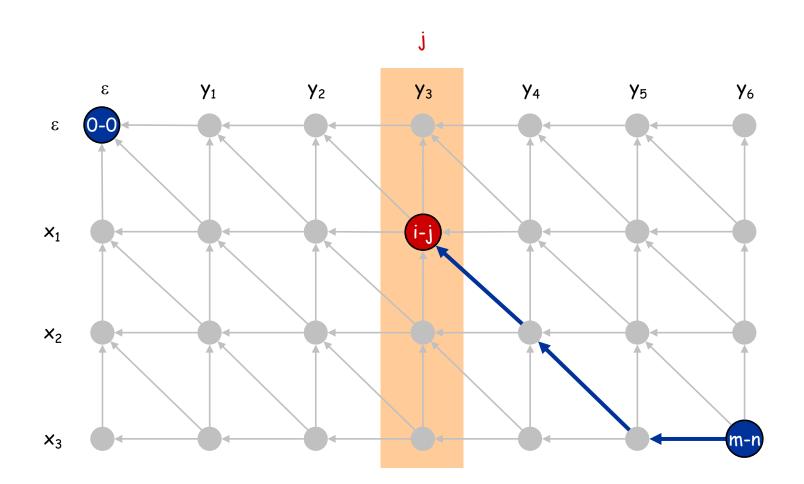
- Let f(i, j) be shortest path from (0,0) to (i, j).
- Can compute $f(\cdot, j)$ for any j in O(mn) time and O(m + n) space.



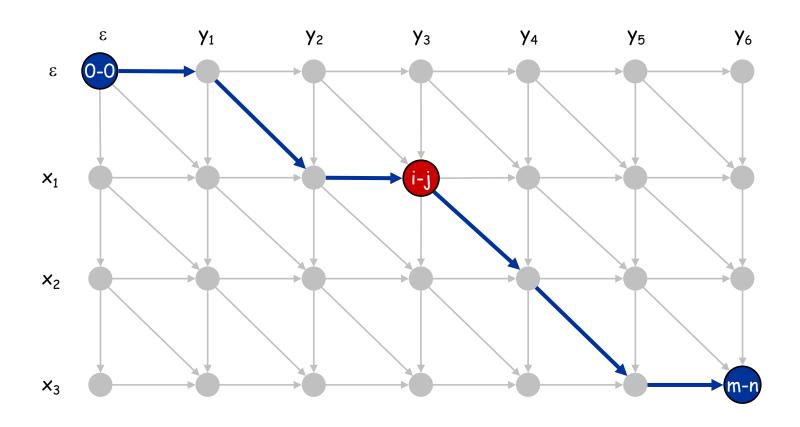
- Let g(i, j) be shortest path from (i, j) to (m, n).
- Can compute $g(\cdot, j)$ by reversing the edge orientations and inverting the roles of (0, 0) and (m, n)



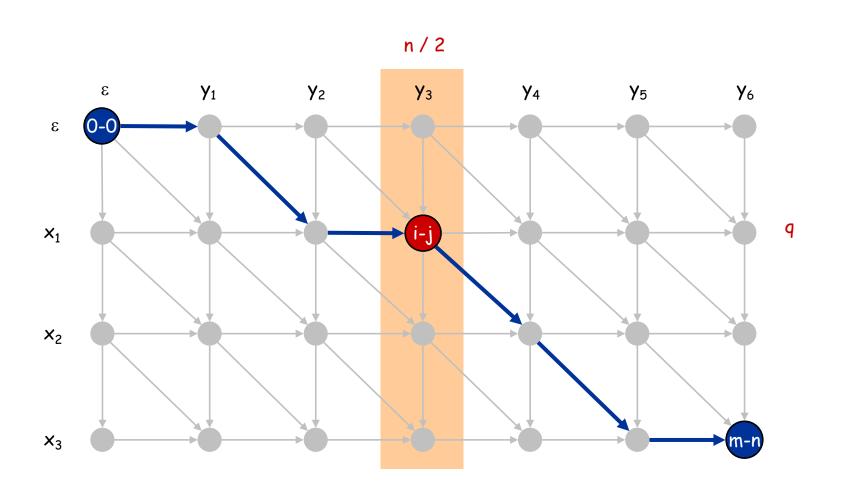
- Let g(i, j) be shortest path from (i, j) to (m, n).
- Can compute $g(\cdot, j)$ for any j in O(mn) time and O(m + n) space.



Observation 1. The cost of the shortest path that uses (i, j) is f(i, j) + g(i, j).



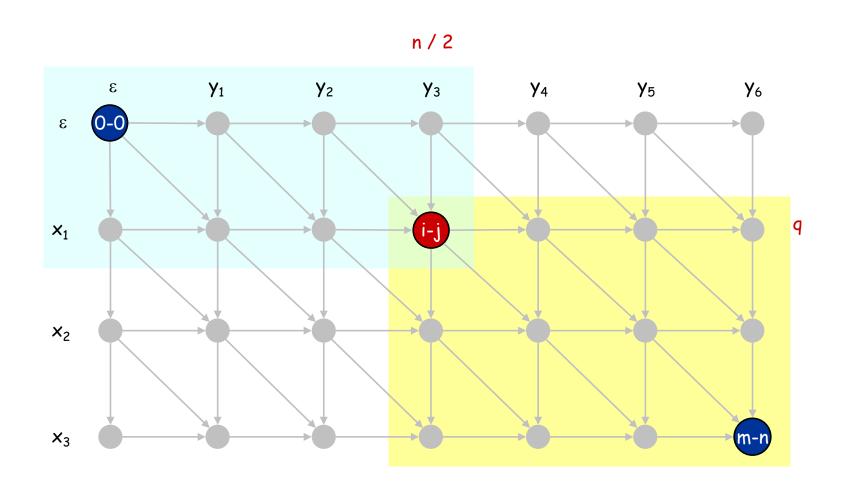
Observation 2. let q be an index that minimizes f(q, n/2) + g(q, n/2). Then, the shortest path from (0, 0) to (m, n) uses (q, n/2).



Divide: find index q that minimizes f(q, n/2) + g(q, n/2) using DP.

• Do alignment at $(x_q, y_{n/2})$.

Conquer: recursively compute optimal alignment in each piece.



Sequence Alignment: Running Time Analysis

Theorem. Let T(m, n) = max running time of algorithm on strings of length m and n. T(m, n) = O(mn).

Pf. (by induction on n)

- O(mn) time to compute $f(\cdot, n/2)$ and $g(\cdot, n/2)$ and find index q.
- T(q, n/2) + T(m q, n/2) time for two recursive calls.
- Choose constant c so that:

$$T(m, 2) \le cm$$

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

- Claim: $T(m, n) \leq 2cmn$
 - Base cases: m = 2 or n = 2.
 - Inductive hypothesis: $T(m', n') \leq 2cm'n'$ with m'<m and n'<n

$$T(m,n) \le T(q,n/2) + T(m-q,n/2) + cmn$$

 $\le 2cqn/2 + 2c(m-q)n/2 + cmn$
 $= cqn + cmn - cqn + cmn$
 $= 2cmn$

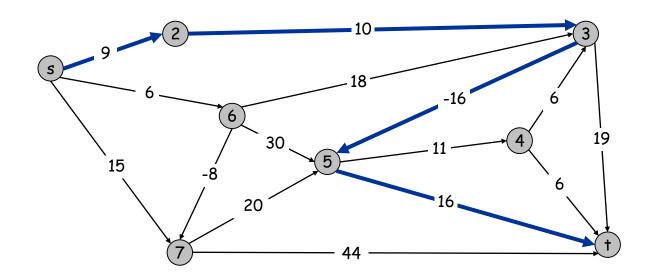
6.8 Shortest Paths

Shortest Paths

Shortest path problem. Given a directed graph G = (V, E), with edge weights c_{vw} , find shortest path from node s to node t.

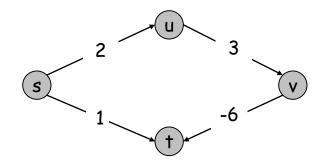
`allow negative weights

Ex. Nodes represent agents in a financial setting and c_{vw} is cost of transaction in which we buy from agent v and sell immediately to w.

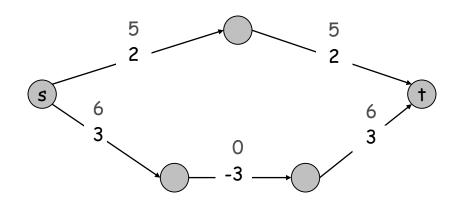


Shortest Paths: Failed Attempts

Dijkstra. Can fail if negative edge costs.

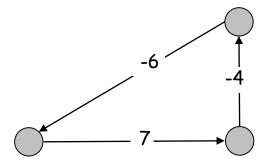


Re-weighting. Adding a constant to every edge weight can fail.

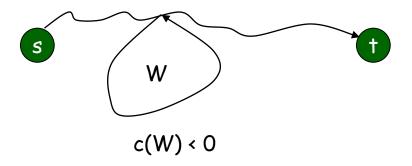


Shortest Paths: Negative Cost Cycles

Negative cost cycle.



Observation. If some path from s to t contains a negative cost cycle, there does not exist a shortest s-t path; otherwise, there exists one that is simple.



Shortest Paths: Dynamic Programming

Def. OPT(i, v) = length of shortest v-t path P using at most i edges.

- Case 1: P uses at most i-1 edges.
 - OPT(i, v) = OPT(i-1, v)
- Case 2: P may use i edges.
 - if (v, w) is first edge, then OPT uses (v, w), and then selects best w-t path using at most i-1 edges

$$OPT(i,v) = \begin{cases} \infty & \text{if } i = 0, v \neq t \\ 0 & \text{if } v = t \\ \min \left\{ OPT(i-1,v), \min_{(v,w) \in E} \left\{ OPT(i-1,w) + c_{vw} \right\} \right\} & \text{otherwise} \end{cases}$$

Remark. By previous observation, if no negative cycles, then OPT(n-1, v) = length of shortest v-t path.

Shortest Paths: Implementation

```
Shortest-Path(G, s, t) {
    foreach node v ∈ V
        M[0, v] ← ∞
    M[0, t] ← 0

for i = 1 to n-1
    foreach node v ∈ V
        M[i, v] ← M[i-1, v]
    foreach edge (v, w) ∈ E
        M[i, v] ← min { M[i, v], M[i-1, w] + c<sub>vw</sub> }

return M[n-1, s]
}
```

Analysis. $\Theta(mn)$ time, $\Theta(n^2)$ space.

Finding the shortest paths. Maintain a "successor" for each table entry.

Shortest Paths: Improvements

```
Shortest-Path(G, s, t) {
    foreach node v ∈ V
        M[0, v] ← ∞
    M[0, t] ← 0

for i = 1 to n-1
    foreach node v ∈ V
        M[i, v] ← M[i-1, v]
    foreach edge (v, w) ∈ E
        M[i, v] ← min { M[i, v], M[i-1, w] + c<sub>vw</sub> }

return M[n-1, s]
}
```

Practical improvements.

• Maintain only one array M[v] = shortest v-t path that we have found so far.

Shortest Paths: Improvements

Practical improvements.

- Maintain only one array M[v] = shortest v-t path that we have found so far.
- No need to check edges of the form (v, w) unless M[w] changed in previous iteration.

Bellman-Ford: Efficient Implementation

```
Push-Based-Shortest-Path(G, s, t) {
   foreach node v \in V
      M[v] \leftarrow \infty
   M[t] \leftarrow 0
   for i = 1 to n-1 {
       foreach node w \in V
          if (M[w] has been updated in previous iteration)
              foreach node v such that (v, w) \in E
                 if (M[v] > M[w] + c_{vw})
                     M[v] \leftarrow M[w] + c_{vw}
       If no M[w] value changed in iteration i, stop.
   return M[s]
```

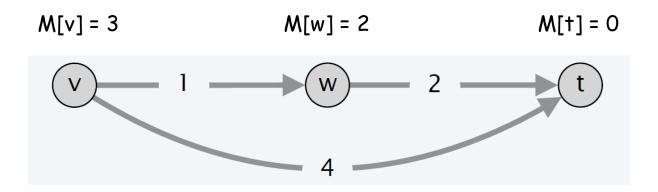
Analysis.

- O(n) extra space
- Time: O(mn) worst case, but substantially faster in practice.

Bellman-Ford: Efficient Implementation

Claim. Throughout the algorithm, M[v] is length of some v-t path, and after i rounds of updates, the value M[v] is the length of shortest v-t path using \leq i edges.

Counter-example:



if nodes w considered before node v, then M[v] = 3 after 1 pass

Bellman-Ford: Efficient Implementation

Claim. Throughout the algorithm, M[v] is length of some v-t path, and after i rounds of updates, the value M[v] is no larger than the length of shortest v-t path using \leq i edges.

6.9 Distance Vector Protocol

Distance Vector Protocol

Communication network.

- Nodes ≈ routers.
- Edges ≈ direct communication link.
- Cost of edge ≈ delay on link. ← naturally nonnegative

Dijkstra's algorithm. Requires global information of network.

Bellman-Ford. Uses only local knowledge of neighboring nodes.

Synchronization. We don't expect routers to run in lockstep. The order in which each foreach loop executes is not important. Moreover, algorithm still converges even if updates are asynchronous.

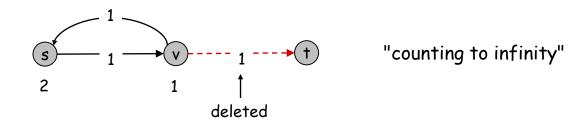
Distance Vector Protocol

Distance vector protocol.

- Each router maintains a vector of shortest path lengths to every other node (distances) and the first hop on each path (directions).
- Algorithm: each router performs n separate computations, one for each potential destination node.
- "Routing by rumor."

Ex. RIP, Xerox XNS RIP, Novell's IPX RIP, Cisco's IGRP, DEC's DNA Phase IV, AppleTalk's RTMP.

Caveat. Edge costs may change during algorithm (or fail completely).



Path Vector Protocols

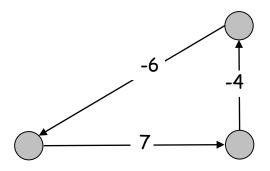
Link state routing.

- not just the distance and first hop
- Each router also stores the entire path.
- Avoids "counting-to-infinity" problem and related difficulties.
- Requires significantly more storage.

Ex. Border Gateway Protocol (BGP), Open Shortest Path First (OSPF).

6.10 Negative Cycles in a Graph

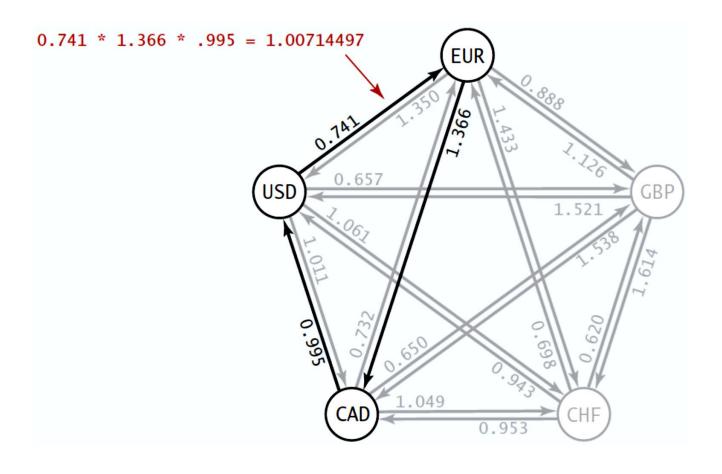
Negative cycle detection problem. Given a digraph G = (V, E), with edge weights c_{vw} , find a negative cycle (if one exists).



Detecting Negative Cycles: Application

Currency conversion. Given n currencies and exchange rates between pairs of currencies, is there an arbitrage opportunity?

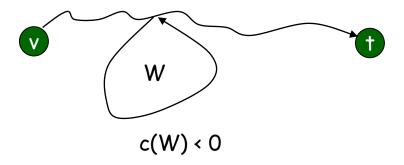
Remark. Fastest algorithm very valuable!



Lemma. If OPT(n,v) = OPT(n-1,v) for all v, then there is no negative cycle with a path to t.

Pf. (by contradiction)

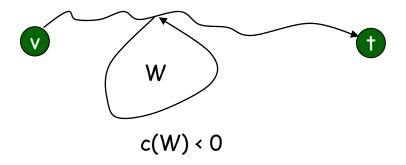
- $OPT(n,v) = OPT(n-1,v) \Rightarrow OPT(i,v) = OPT(n-1,v)$ for $i \ge n$
- But negative cycle in a path implies that OPT(i,v) always decreases as i increases



Lemma. If OPT(n,v) < OPT(n-1,v) for some node v, then (any) shortest path from v to t contains a cycle W. Moreover W has negative cost.

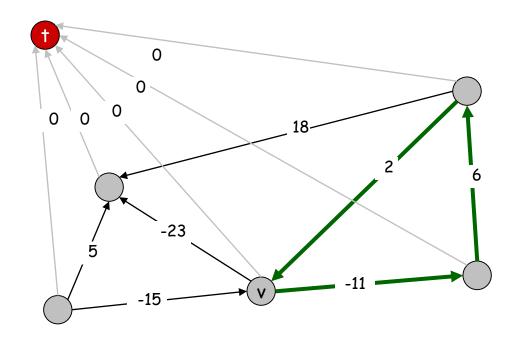
Pf. (by contradiction)

- Since OPT(n,v) < OPT(n-1,v), we know the shortest v-t path P has exactly n edges.
- By pigeonhole principle, P must contain a directed cycle W.
- Deleting W yields a v-t path with < n edges \Rightarrow W has negative cost.



Theorem. Can detect negative cost cycle in O(mn) time.

- Add new node t and connect all nodes to t with 0-cost edge.
- Check if OPT(n, v) = OPT(n-1, v) for all nodes v.
 - if yes, then no negative cycles
 - if no, then extract cycle from shortest path from v to t



Dynamic Programming: Chapter Summary

Dynamic Programming

Basic idea

- Polynomial number of sub-problems with a natural ordering from smallest to largest.
- Optimal solution to a sub-problem can be constructed from optimal solutions of smaller sub-problems.
- Sub-problems are overlapping!

Guideline

- Define the sub-problems
 - OPT(...)
- Write down the recursive formulas
 - Ex: OPT(i) = max(f(OPT(j)), g(OPT(k)), ...), j,k < i
- Compute the formulas either bottom-up or top-down

Dynamic Programming

Algorithms

- Weighted interval scheduling
 - 1D array; binary choice
- Knapsack
 - 2D array; adding a new variable (weight limit)
- RNA secondary structure
 - 2D array: intervals
- Sequence Alignment
 - 2D array: prefix alignment
- Sequence Alignment in Linear Space
 - Combination of divide-and-conquer and dynamic programming
- Shortest path with negative edges
 - (Bellman-Ford) 2D array: shortest path with edge number ≤ i
- Distance Vector Protocol
- Negative Cycle Detection