Dynamic Programming

T. M. Murali

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4. Dynamic programming
   - More powerful than greedy and divide-and-conquer strategies.
   - *Implicitly* explore space of all possible solutions.
   - Solve multiple sub-problems and build up correct solutions to larger and larger sub-problems.
   - Careful analysis needed to ensure number of sub-problems solved is polynomial in the size of the input.

History of Dynamic Programming

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- The Secretary of Defense at that time was hostile to mathematical research.
- Bellman sought an impressive name to avoid confrontation.
  - "it's impossible to use dynamic in a pejorative sense"
  - "something not even a Congressman could object to" (Bellman, R. E., *Eye of the Hurricane, An Autobiography*).

Applications of Dynamic Programming

- Computational biology: Smith-Waterman algorithm for sequence alignment.
- Operations research: Bellman-Ford algorithm for shortest path routing in networks.
- Control theory: Viterbi algorithm for hidden Markov models.
- Computer science (theory, graphics, AI, ...): Unix `diff` command for comparing two files.

Review: Interval Scheduling

INTERVAL SCHEDULING

INSTANCE: Nonempty set $\{(s_i, f_i), 1 \leq i \leq n\}$ of start and finish times of n jobs.

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SOLUTION: The largest subset of mutually compatible jobs.

- Two jobs are *compatible* if they do not overlap.
- Greedy algorithm: sort jobs in increasing order of finish times. Add next job to current subset only if it is compatible with previously-selected jobs.

Weighted Interval Scheduling

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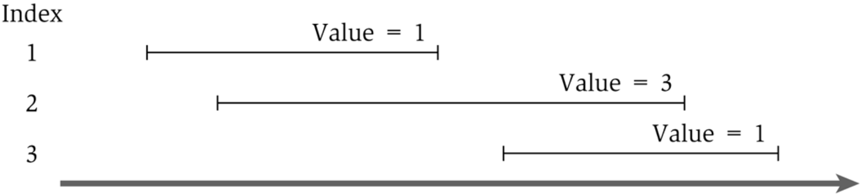
INSTANCE: Nonempty set $\{(s_i, f_i), 1 \leq i \leq n\}$ of start and finish times of n jobs and a weight $v_i \geq 0$ associated with each job.

SOLUTION: A set S of mutually compatible jobs such that $\sum_{i \in S} v_i$ is maximised.



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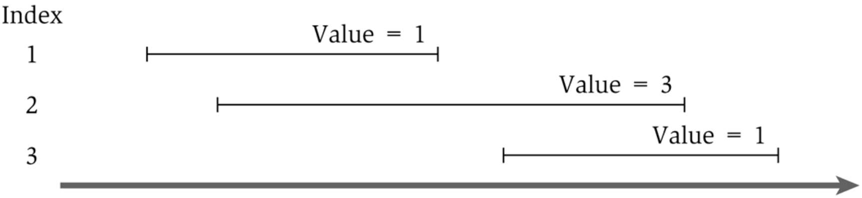

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- Greedy algorithm can produce arbitrarily bad results for this problem.

Approach

- Sort jobs in increasing order of finish time and relabel: $f_1 \leq f_2 \leq \ldots \leq f_n$.
- Request i comes before request j if $i < j$.
- $p(j)$ is the largest index $i < j$ such that job i is compatible with job j. $p(j) = 0$ if there is no such job i.

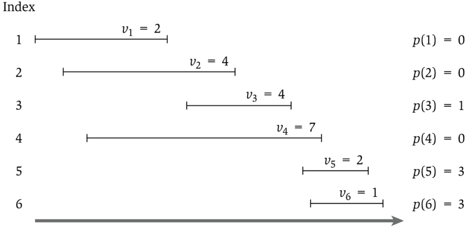
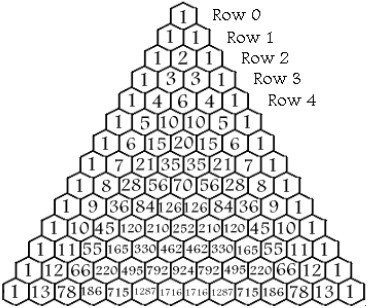


Figure 6.2 An instance of weighted interval scheduling with the functions $p(j)$ defined for each interval j.

- We will develop optimal algorithm from obvious statements about the problem.

Detour: a Binomial Identity



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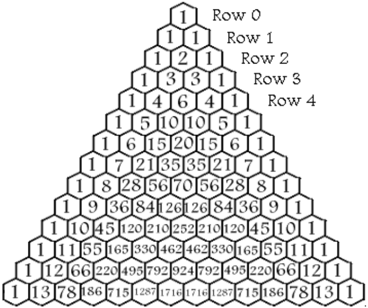

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  - Each element is a binomial co-efficient.
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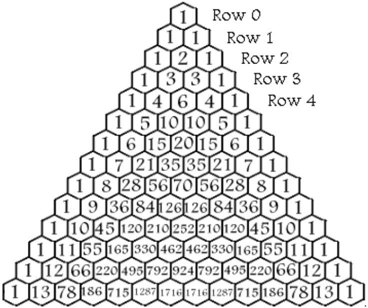

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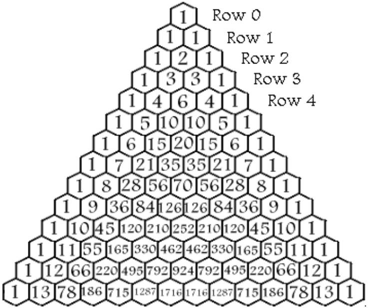

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- Proof: either we select the nth element or not ...

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- $\mathcal{O}$ must be the best of these two choices!
- Suggests finding optimal solution for sub-problems consisting of jobs $\{1, 2, \ldots, j-1, j\}$, for all values of j.

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- When does request j belong to $\mathcal{O}_j$? If and only if $v_j + \text{OPT}(p(j)) \geq \text{OPT}(j-1)$.

Recursive Algorithm

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Compute-Opt(j)
  If j = 0 then
    Return 0
  Else
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- What is the running time of the algorithm? Can be exponential in n.
- When $p(j) = j - 2$, for all $j \geq 2$: recursive calls are for $j - 1$ and $j - 2$.

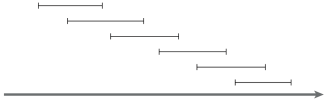

Figure 6.4 An instance of weighted interval scheduling on which the simple Compute-Opt recursion will take exponential time. The values of all intervals in this instance are 1.

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M-Compute-Opt(j)
  If j = 0 then
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  Else if M[j] is not empty then
    Return M[j]
  Else
    Define M[j] = max(v_j + M-Compute-Opt(p(j)), M-Compute-Opt(j − 1))
    Return M[j]
  Endif
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- How many such recursive calls are there in total?
- Use number of filled entries in M as a measure of progress.
- Each time `M-Compute-Opt` issues two recursive calls, it fills in a new entry in M.
- Therefore, total number of recursive calls is $O(n)$.

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- Recall: request j belong to $\mathcal{O}_j$ if and only if $v_j + \text{OPT}(p(j)) \geq \text{OPT}(j-1)$.
- Can recover $\mathcal{O}_j$ from values of the optimal solutions in $O(j)$ time.

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- Can recover $\mathcal{O}_j$ from values of the optimal solutions in $O(j)$ time.

```
Find-Solution(j)
  If j = 0 then
    Output nothing
  Else
    If v_j + M[p(j)] ≥ M[j − 1] then
      Output j together with the result of Find-Solution(p(j))
    Else
      Output the result of Find-Solution(j − 1)
    Endif
  Endif
```

From Recursion to Iteration

- Unwind the recursion and convert it into iteration.
- Can compute values in M iteratively in $O(n)$ time.
- `Find-Solution` works as before.

```
Iterative-Compute-Opt
  M[0] = 0
  For j = 1, 2, ..., n
    M[j] = max(v_j + M[p(j)], M[j − 1])
  Endfor
```

Basic Outline of Dynamic Programming

- To solve a problem, we need a collection of sub-problems that satisfy a few properties:
  1. There are a polynomial number of sub-problems.
  2. The solution to the problem can be computed easily from the solutions to the sub-problems.
  3. There is a natural ordering of the sub-problems from “smallest” to “largest”.
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  4. There is an easy-to-compute recurrence that allows us to compute the solution to a sub-problem from the solutions to some smaller sub-problems.
- Difficulties in designing dynamic programming algorithms:
  1. Which sub-problems to define?
  2. How can we tie together sub-problems using a recurrence?
  3. How do we order the sub-problems (to allow iterative computation of optimal solutions to sub-problems)?

Least Squares Problem

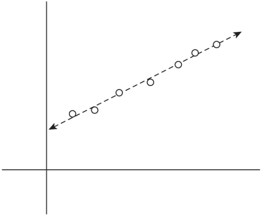

Figure 6.6 A "line of best fit."

- Given scientific or statistical data plotted on two axes.
- Find the "best" line that "passes" through these points.

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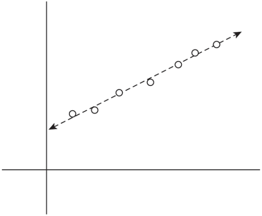

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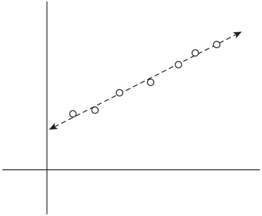

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INSTANCE: Set $P = \{(x_1, y_1), (x_2, y_2), \ldots, (x_n, y_n)\}$ of n points.
SOLUTION: Line $L : y = ax + b$ that minimises

$$\text{Error}(L, P) = \sum_{i=1}^{n} (y_i - ax_i - b)^2.$$

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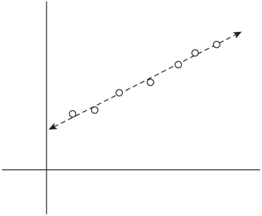

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- Solution is achieved by

$$a = \frac{n \sum_i x_i y_i - \left(\sum_i x_i\right)\left(\sum_i y_i\right)}{n \sum_i x_i^2 - \left(\sum_i x_i\right)^2} \text{ and } b = \frac{\sum_i y_i - a \sum_i x_i}{n}$$

Segmented Least Squares

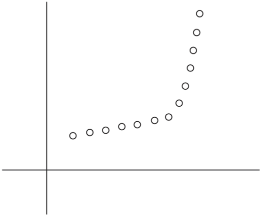

Figure 6.7 A set of points that lie approximately on two lines.

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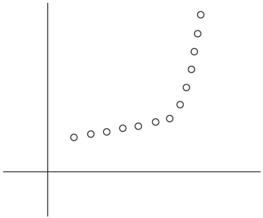

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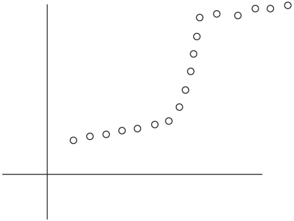

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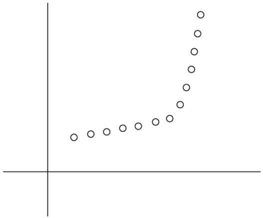

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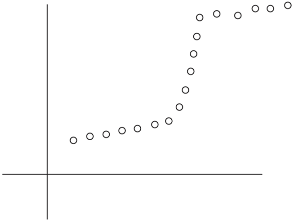

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- Want to fit multiple lines through P.
- Each line must fit contiguous set of x-coordinates.
- Lines must minimise total error.

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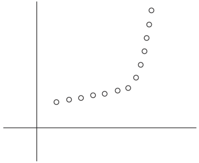

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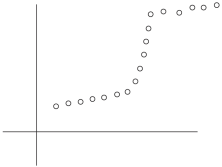
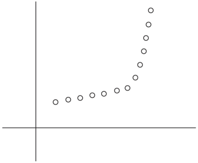


Figure 6.8 A set of points that lie approximately on three lines.

Segmented Least Squares



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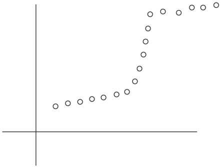

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SEGMENTED LEAST SQUARES

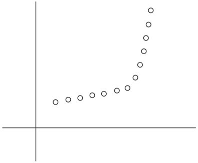
INSTANCE: Set $P = \{p_i = (x_i, y_i), 1 \leq i \leq n\}$ of n points, $x_1 < x_2 < \cdots < x_n$.

SOLUTION: A integer k, a partition of P into k segments $\{P_1, P_2, \ldots, P_k\}$, k lines $L_j : y = a_j x + b_j, 1 \leq j \leq k$ that minimise

$$\sum_{j=1}^{k} \text{Error}(L_j, P_j)$$

- A subset P' of P is a *segment* if $1 \leq i < j \leq n$ exist such that $P' = \{(x_i, y_i), (x_{i+1}, y_{i+1}), \ldots, (x_{j-1}, y_{j-1}), (x_j, y_j)\}$.

Segmented Least Squares



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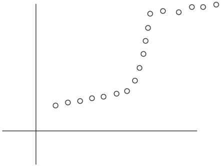

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SEGMENTED LEAST SQUARES

INSTANCE: Set $P = \{p_i = (x_i, y_i), 1 \le i \le n\}$ of n points, $x_1 < x_2 < \cdots < x_n$ and a parameter $C > 0$.

SOLUTION: A integer k, a partition of P into k segments $\{P_1, P_2, \ldots, P_k\}$, k lines $L_j : y = a_j x + b_j, 1 \le j \le k$ that minimise

$$\sum_{j=1}^{k} \mathrm{Error}(L_j, P_j) + Ck.$$

- A subset P' of P is a *segment* if $1 \le i < j \le n$ exist such that $P' = \{(x_i, y_i), (x_{i+1}, y_{i+1}), \ldots, (x_{j-1}, y_{j-1}), (x_j, y_j)\}$.

Formulating the Recursion I

- Observation: p_n is part of some segment in the optimal solution. This segment starts at some point p_i.
- Let $OPT(i)$ be the optimal value for the points $\{p_1, p_2, \ldots, p_i\}$.
- Let $e_{i,j}$ denote the minimum error of any line that fits $\{p_i, p_2, \ldots, p_j\}$.
- We want to compute $\mathrm{OPT}(n)$.

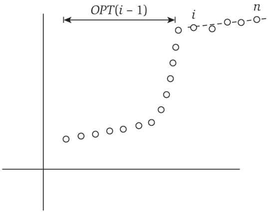

Figure 6.9 A possible solution: a single line segment fits points $p_i, p_{i+1}, \ldots, p_n$, and then an optimal solution is found for the remaining points $p_1, p_2, \ldots, p_{i-1}$.

- If the last segment in the optimal partition is $\{p_i, p_{i+1}, \ldots, p_n\}$, then

$$\mathrm{OPT}(n) = e_{i,n} + C + \mathrm{OPT}(i-1)$$

Formulating the Recursion II

- Consider the sub-problem on the points $\{p_1, p_2, \dots p_j\}$
- To obtain $\text{OPT}(j)$, if the last segment in the optimal partition is $\{p_i, p_{i+1}, \dots, p_j\}$, then

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- Since i can take only j distinct values,

$$\mathrm{OPT}(j) = \min_{1 \leq i \leq j} \left(e_{i,j} + C + \mathrm{OPT}(i-1) \right)$$

- Segment $\{p_i, p_{i+1}, \ldots p_j\}$ is part of the optimal solution for this sub-problem if and only if the minimum value of $\mathrm{OPT}(j)$ is obtained using index i.

Dynamic Programming Algorithm

$$\mathrm{OPT}(j) = \min_{1 \leq i \leq j} \left(e_{i,j} + C + \mathrm{OPT}(i-1) \right)$$

```
Segmented-Least-Squares(n)
  Array M[0...n]
  Set M[0]=0
  For all pairs i≤j
    Compute the least squares error e_{i,j} for the segment p_i,...,p_j
  Endfor
  For j=1,2,...,n
    Use the recurrence (6.7) to compute M[j]
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  Endfor
  Return M[n]
```

- Running time is $O(n^3)$, can be improved to $O(n^2)$.
- We can find the segments in the optimal solution by backtracking.

RNA Molecules

- RNA is a basic biological molecule. It is single stranded.
- RNA molecules fold into complex "secondary structures."
- Secondary structure often governs the behaviour of an RNA molecule.
- Various rules govern secondary structure formation:

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1. Pairs of bases match up; each base matches with ≤ 1 other base.
2. Adenine always matches with Uracil.
3. Cytosine always matches with Guanine.
4. There are no kinks in the folded molecule.
5. Structures are "knot-free".

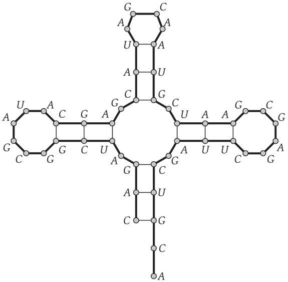

Figure 6.13 An RNA secondary structure. Thick lines connect adjacent elements of the sequence; thin lines indicate pairs of elements that are matched.

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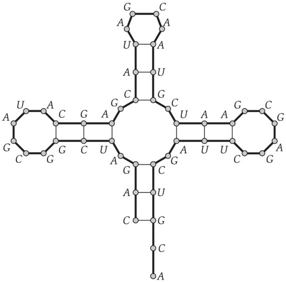

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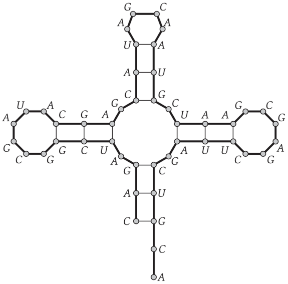

Figure 6.13 An RNA secondary structure. Thick lines connect adjacent elements of the sequence; thin lines indicate pairs of elements that are matched.

- Problem: given an RNA molecule, predict its secondary structure.
- Hypothesis: In the cell, RNA molecules form the secondary structure with the lowest total free energy.

Formulating the Problem

- An *RNA molecule* is a string $B = b_1 b_2 \ldots b_n$; each $b_i \in \{A, C, G, U\}$.
- A *secondary structure on B* is a set of pairs $S = \{(i, j)\}$, where $1 \leq i, j \leq n$ and

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- A *secondary structure on B* is a set of pairs $S = \{(i, j)\}$, where $1 \leq i, j \leq n$ and
  1. *(No kinks.)* If $(i, j) \in S$, then $i < j - 4$.
  2. *(Watson-Crick)* The elements in each pair in S consist of either $\{A, U\}$ or $\{C, G\}$ (in either order).
  3. S is a *matching*: no index appears in more than one pair.
  4. *(No knots)* If (i, j) and (k, l) are two pairs in S, then we cannot have $i < k < j < l$.

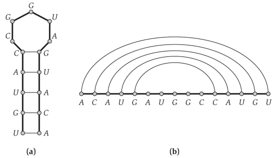

Figure 6.14 Two views of an RNA secondary structure. In the second view, (b), the string has been "stretched" lengthwise, and edges connecting matched pairs appear as noncrossing "bubbles" over the string.

- The *energy* of a secondary structure $\propto$ the number of base pairs in it.

Dynamic Programming Approach

- $OPT(j)$ is the maximum number of base pairs in a secondary structure for $b_1 b_2 \ldots b_j$.

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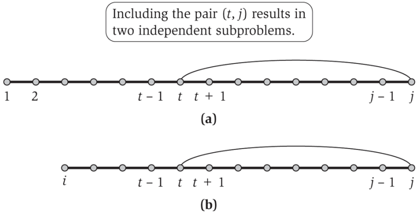

Figure 6.15 Schematic views of the dynamic programming recurrence using (a) one variable, and (b) two variables.

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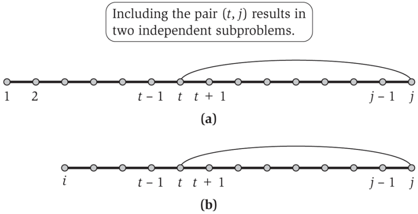

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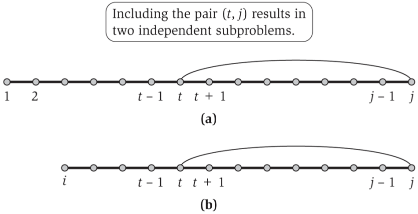
- Insight: need sub-problems indexed both by start and by end.



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- In the "inner" maximisation, t runs over all indices between i and $j-5$ that are allowed to pair with j.

Dynamic Programming Algorithm

$$\mathrm{OPT}(i,j) = \max\left(\mathrm{OPT}(i,j-1), \max_t \left(1 + \mathrm{OPT}(i,t-1) + \mathrm{OPT}(t+1,j-1)\right)\right)$$

- There are $O(n^2)$ sub-problems.
- How do we order them from "smallest" to "largest"?

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Initialize OPT(i, j) = 0 whenever i ≥ j − 4
For k = 5, 6, . . . , n − 1
  For i = 1, 2, . . . n − k
    Set j = i + k
    Compute OPT(i, j) using the recurrence in (6.13)
  Endfor
Endfor
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```

- Running time of the algorithm is $O(n^3)$.

Example of Algorithm

RNA sequence *ACCGGUAGU*

4	0	0	0	
3	0	0		
2	0			
$i = 1$				
	$j = 6$	7	8	9

Initial values

4	0	0	0	0
3	0	0	1	
2	0	0		
$i = 1$	1			
	$j = 6$	7	8	9

Filling in the values for $k = 5$

4	0	0	0	0
3	0	0	1	1
2	0	0	1	
$i = 1$	1	1		
	$j = 6$	7	8	9

Filling in the values for $k = 6$

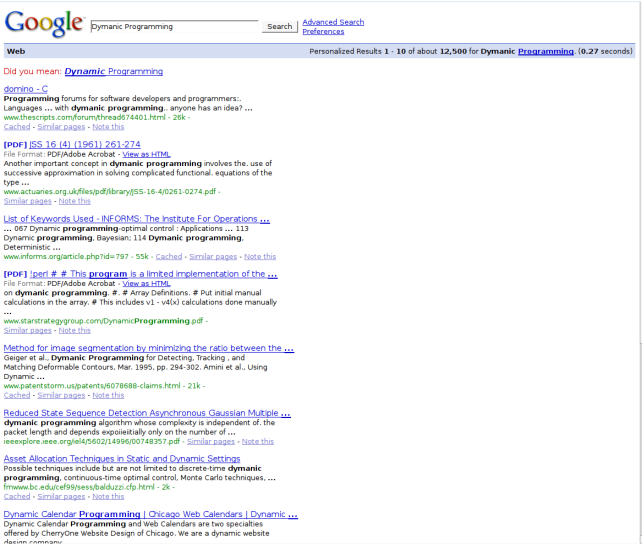
4	0	0	0	0
3	0	0	1	1
2	0	0	1	1
$i = 1$	1	1	1	
	$j = 6$	7	8	9

Filling in the values for $k = 7$

4	0	0	0	0
3	0	0	1	1
2	0	0	1	1
$i = 1$	1	1	1	2
	$j = 6$	7	8	9

Filling in the values for $k = 8$

Google Search for "Dymanic Programming"



- How do they know "Dynamic" and "Dymanic" are similar?

Sequence Similarity

- Given two strings, measure how similar they are.
- Given a database of strings and a query string, compute the string most similar to query in the database.
- Applications:
  - Online searches (Web, dictionary).
  - Spell-checkers.
  - Computational biology
  - Speech recognition.
  - Basis for Unix `diff`.

Defining Sequence Similarity

```
o-currance
occurrence
```

```
o-curr-ance
occurre-nce
```

```
abbbaa--bbbbaab
ababaaabbbba-b
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```

- *Edit distance* model: how many changes must you to make to one string to transform it into another?
- Changes allowed are deleting a letter, adding a letter, changing a letter.

Edit Distance

```
o-currance      o-curr-ance
occurrence      occurre-nce
```

- Proposed by Needleman and Wunsch in the early 1970s.
- Input: two strings $x = x_1 x_2 x_3 \ldots x_m$ and $y = y_1 y_2 \ldots y_n$.
- Sets $\{1, 2, \ldots, m\}$ and $\{1, 2, \ldots, n\}$ represent positions in x and y.

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o-currance                                    o-curr-ance
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- A *matching* of these sets is a set M of ordered pairs such that
  1. in each pair (i, j), $1 \leq i \leq m$ and $1 \leq j \leq n$ and
  2. no index from x (respectively, from y) appears as the first (respectively, second) element in more than one ordered pair.
- An index is *not matched* if it does not appear in the matching.

Edit Distance

```
o-currance          o-curr-ance
occurrence          occurre-nce
```

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- Input: two strings $x = x_1x_2x_3 \dots x_m$ and $y = y_1y_2 \dots y_n$.
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- An index is *not matched* if it does not appear in the matching.
- A matching M is an *alignment* if there are no "crossing pairs" in M: if $(i, j) \in M$ and $(i', j') \in M$ and $i < i'$ then $j < j'$.

Edit Distance

```
o-currance          o-curr-ance
occurrence          occurre-nce
```

- Proposed by Needleman and Wunsch in the early 1970s.
- Input: two strings $x = x_1 x_2 x_3 \dots x_m$ and $y = y_1 y_2 \dots y_n$.
- Sets $\{1, 2, \dots, m\}$ and $\{1, 2, \dots, n\}$ represent positions in x and y.
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- Cost of an alignment is the sum of gap and mismatch penalties:

  Gap penalty Penalty $\delta > 0$ for every unmatched index.

  Mismatch penalty Penalty $\alpha_{x_i y_j} > 0$ if $(i, j) \in M$ and $x_i \neq y_j$.

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  Mismatch penalty: Penalty $\alpha_{x_i y_j} > 0$ if $(i, j) \in M$ and $x_i \neq y_j$.
- Output: compute an alignment of minimal cost.

Dynamic Programming Approach

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

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  - $(i, j) \in M$ if and only if minimum is achieved by the first term.
- What are the base cases? $\text{OPT}(i, 0) = \text{OPT}(0, i) = i\delta$.

Dynamic Programming Algorithm

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Alignment(X,Y)
  Array A[0...m, 0...n]
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         Use the recurrence (6.16) to compute A[i,j]
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- Running time is $O(mn)$. Space used in $O(mn)$.

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- Running time is $O(mn)$. Space used in $O(mn)$.
- Can compute $\mathrm{OPT}(m,n)$ in $O(mn)$ time and $O(m+n)$ space (Hirschberg 1975, Chapter 6.7).
- Can compute *alignment* in the same bounds by combining dynamic programming with divide and conquer.

Graph-theoretic View of Sequence Alignment

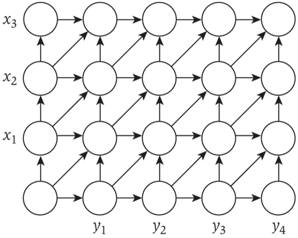

Figure 6.17 A graph-based picture of sequence alignment.

- Grid graph G_{xy}:
  - Rows labelled by symbols in x and columns labelled by symbols in y.
  - Edges from node (i, j) to $(i, j+1)$), to $(i+1, j)$, and to $(i+1, j+1)$.
  - Edges directed upward and to the right have cost δ.
  - Edge directed from (i, j) to $(i+1, j+1)$ has cost $\alpha_{x_{i+1} y_{j+1}}$.

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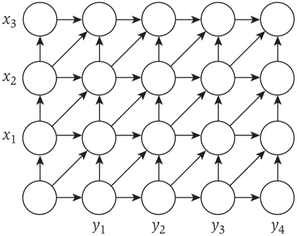

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  - Edges directed upward and to the right have cost δ.
  - Edge directed from (i, j) to $(i + 1, j + 1)$ has cost $\alpha_{x_{i+1} y_{j+1}}$.
- *f(i, j)*: minimum cost of a path in G_{XY} from $(0, 0)$ to (i, j).
- Claim: $f(i, j) = \text{OPT}(i, j)$ and diagonal edges in the shortest path are the matched pairs in the alignment.

Motivation

- Computational finance:
  - Each node is a financial agent.
  - The cost c_{uv} of an edge (u, v) is the cost of a transaction in which we buy from agent u and sell to agent v.
  - Negative cost corresponds to a profit.
- Internet routing protocols
  - Dijkstra's algorithm needs knowledge of the entire network.
  - Routers only know which other routers they are connected to.
  - Algorithm for shortest paths with negative edges is decentralised.
  - We will not study this algorithm in the class. See Chapter 6.9.

Problem Statement

- Input: a directed graph $G = (V, E)$ with a cost function $c : E \rightarrow \mathbb{R}$, i.e., c_{uv} is the cost of the edge $(u, v) \in E$.
- A *negative cycle* is a directed cycle whose edges have a total cost that is negative.
- Two related problems:
  1. If G has no negative cycles, find the *shortest s-t path*: a path of from source s to destination t with minimum total cost.
  2. Does G have a *negative cycle*?

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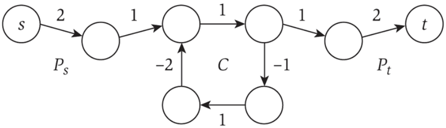

Figure 6.20 In this graph, one can find *s*-*t* paths of arbitrarily negative cost (by going around the cycle *C* many times).

Approaches for Shortest Path Algorithm

1. Dijsktra's algorithm.

2. Add some large constant to each edge.

Approaches for Shortest Path Algorithm

1. Dijsktra's algorithm. Computes incorrect answers because it is greedy.
2. Add some large constant to each edge. Computes incorrect answers because the minimum cost path changes.

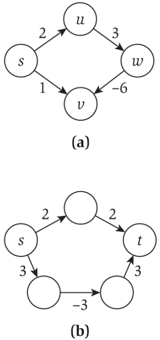

Figure 6.21 (a) With negative edge costs, Dijkstra's Algorithm can give the wrong answer for the Shortest-Path Problem. (b) Adding 3 to the cost of each edge will make all edges nonnegative, but it will change the identity of the shortest s-t path.

Dynamic Programming Approach

- Assume G has no negative cycles.
- Claim: There is a shortest path from s to t that is *simple* (does not repeat a node)

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  - Shortest s-t path has $\leq n - 1$ edges: how we can reach t using i edges, for different values of i?
  - We do not know which nodes will be in shortest s-t path: how we can reach t from each node in V?

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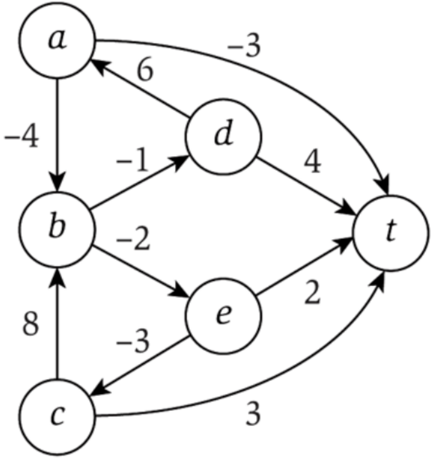
  - We do not know which nodes will be in shortest s-t path: how we can reach t from each node in V?
- Sub-problems defined by varying the number of edges in the shortest path and by varying the starting node in the shortest path.



Dynamic Programming Recursion

- $OPT(i, v)$: minimum cost of a v-t path that uses at most i edges.
- t is not explicitly mentioned in the sub-problems.
- Goal is to compute $\text{OPT}(n-1, s)$.

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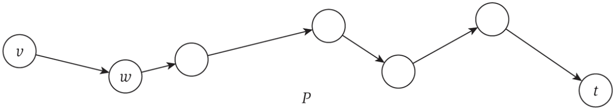

Figure 6.22 The minimum-cost path P from v to t using at most i edges.

- Let P be the optimal path whose cost is $\mathrm{OPT}(i, v)$.

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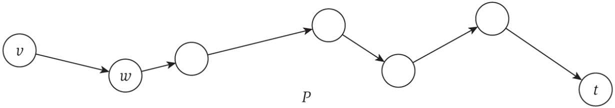

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- Let P be the optimal path whose cost is $\text{OPT}(i, v)$.
    1. If P actually uses $i-1$ edges, then $\text{OPT}(i, v) = \text{OPT}(i-1, v)$.
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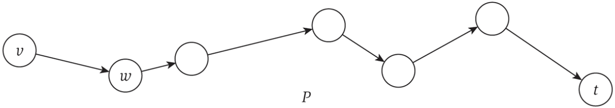

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Alternate Dynamic Programming Formulation

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$$\mathrm{OPT}(i, v) = \min_{w \in V} \left(c_{vw} + \mathrm{OPT}(i-1, w) \right)$$

- Compare the recurrence above to the previous recurrence:

$$\mathrm{OPT}(i, v) = \min \left(\mathrm{OPT}(i-1, v), \min_{w \in V} \left(c_{vw} + \mathrm{OPT}(i-1, w) \right) \right)$$

Bellman-Ford Algorithm

$$\mathrm{OPT}(i, v) = \min\left(\mathrm{OPT}(i-1, v), \min_{w \in V}\left(c_{vw} + \mathrm{OPT}(i-1, w)\right)\right)$$

```
Shortest-Path(G, s, t)
  n = number of nodes in G
  Array M[0 ... n − 1, V]
  Define M[0, t] = 0 and M[0, v] = ∞ for all other v ∈ V
  For i = 1, ..., n − 1
    For v ∈ V in any order
      Compute M[i, v] using the recurrence (6.23)
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```

- Space used is $O(n^2)$. Running time is $O(n^3)$.
- If shortest path uses k edges, we can recover it in $O(kn)$ time by tracing back through smaller sub-problems.

An Improved Bound on the Running Time

- Suppose G has n nodes and $m \ll \binom{n}{2}$ edges. Can we demonstrate a better upper bound on the running time?

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- If n_v is the number of neighbours of v, then in each round, we spend time equal to

$$\sum_{v \in V} n_v = m.$$

- The total running time is $O(mn)$.

Improving the Memory Requirements

$$M[i, v] = \min\left(M[i-1, v], \min_{w \in V}\left(c_{vw} + M[i-1, w]\right)\right)$$

- The algorithm uses $O(n^2)$ space to store the array M.

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- Observe that $M[i, v]$ depends only on $M[i-1, *]$ and no other indices.

Improving the Memory Requirements

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  1. Maintain two arrays M and N indexed over V.
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- Claim: at the beginning of iteration i, M stores values of $\text{OPT}(i-1, v)$ for all nodes $v \in V$.
- Space used is $O(n)$.

Computing the Shortest Path: Algorithm

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- How can we recover the shortest path that has cost $M[v]$?
- For each node v, maintain $f(v)$, the first node after v in the current shortest path from v to t.
- To maintain $f(v)$, if we ever set $M[v]$ to $\min_{w \in V}\left(c_{vw} + N[w]\right)$, set $f(v)$ to be the node w that attains this minimum.
- At the end, follow $f(v)$ pointers from s to t.

Computing the Shortest Path: Correctness

- *Pointer graph* $P(V, F)$: each edge in F is $(v, f(v))$.
  - Can P have cycles?
  - Is there a path from s to t in P?
  - Can there be multiple paths s to t in P?
  - Which of these is the shortest path?

Computing the Shortest Path: Cycles in P

$$M[v] = \min\left(N[v], \min_{w \in V}\left(c_{vw} + N[w]\right)\right)$$

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  - What is the situation just before this addition?

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  - Adding all these inequalities, $0 > \sum_{i=1}^{k-1} c_{v_i v_{i+1}} + c_{v_k v_1} =$ cost of C.

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  - Adding all these inequalities, $0 > \sum_{i=1}^{k-1} c_{v_i v_{i+1}} + c_{v_k v_1} =$ cost of C.
- Corollary: if G has no negative cycles that P does not either.

Computing the Shortest Path: Paths in P

- Let P be the pointer graph upon termination of the algorithm.
- Consider the path P_v in P obtained by following the pointers from v to $f(v) = v_1$, to $f(v_1) = v_2$, and so on.

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- Claim: P_v terminates at t.
- Claim: P_v is the shortest path in G from v to t.

Bellman-Ford Algorithm: Early Termination

$$M[v] = \min\left(N[v], \min_{w \in V}\left(c_{vw} + N[w]\right)\right)$$

- In general, after i iterations, the path whose length is $M[v]$ may have many more than i edges.

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- In general, after i iterations, the path whose length is $M[v]$ may have many more than i edges.
- Early termination: If M equals N after processing all the nodes, we have computed all the shortest paths to t.