Dynamic Programming*

Xiaofeng Gao

Department of Computer Science and Engineering Shanghai Jiao Tong University, P.R.China

Algorithm Course: Shanghai Jiao Tong University

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Basic Methodology

our Examples (Recipe of Dynamic Programn

Algorithmic Paradigms

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

Divide-and-conquer: Break up a problem into sub-problems, solve each sub-problem independently, 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.

Basic Methodolo A Comprehensive Exam

Outline

- Basic Methodology
 - Introduction
 - Four Examples (Recipe of Dynamic Programming)
- 2 A Comprehensive Example
 - String Similarity
 - Sequence Alignment in Linear Space

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Basic Methodology A Comprehensive Example

Introduction
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Dynamic Programming Applications

Areas

- Bioinformatics.
- o Control theory.
- o Information theory.
- o Operations research.
- o Computer science: theory, graphics, AI, compilers, systems, ...

Some famous dynamic programming algorithms

- Unix diff for comparing two files.
- Viterbi for hidden Markov models.
- o Smith-Waterman for genetic sequence alignment.
- Bellman-Ford for shortest path routing in networks.
- o Cocke-Kasami-Younger for parsing context free grammars.

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Four Examples (Recipe of Dynamic Programming)

Four Examples

- Weighted Interval Scheduling
- Segmented Least Squares
- Knapsack Problem
- RNA Secondary Structure

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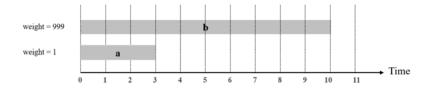
Four Examples (Recipe of Dynamic Programming)

Unweighted Interval Scheduling Review

Recall: Greedy algorithm works if all weights are 1.

- o 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.



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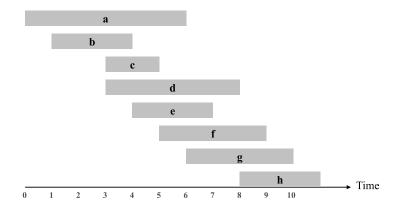
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Weighted Interval Scheduling Problem

Job j starts at s_i , finishes at f_i , and has weight or value v_i .

Two jobs compatible if they don't overlap.

Goal: find maximum weight subset of mutually compatible jobs.



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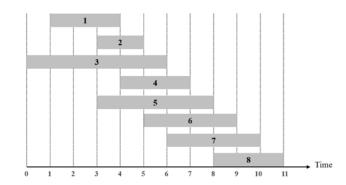
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Weighted Interval Scheduling

Notation: Label jobs by finishing time: $f_1 \le f_2 \le \cdots \le f_n$.

Definition: p(j) = largest index i < j such that job i is compatible with j.

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



Binary Choice

Greedy template: OPT(j) = value of optimal solution to the problem consisting of job requests $1, 2, \dots, j$.

Optimal substructure:

Case 1: OPT selects job j.

- \circ collect profit v_i
- can't use incompatible jobs $\{p(j) + 1, p(j) + 2, \dots, j 1\}$
- must include optimal solution to problem consisting of remaining compatible jobs $1, 2, \dots, p(j)$

Case 2: OPT does not select job *j*.

• must include optimal solution to problem consisting of remaining compatible jobs $1, 2, \dots, j-1$

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

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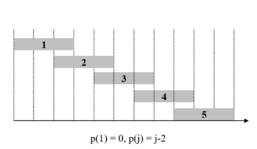
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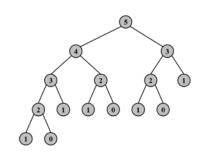
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Brute Force Algorithm

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

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





Brute Force Algorithm

```
Algorithm 1: Brute Force
Input: n; s_1, \dots, s_n; f_1, \dots, f_n; v_1, \dots, v_n;

1 Sort jobs by finish times so that f_1 \le f_2 \le \dots \le f_n;

2 Compute p(1), p(2), \dots, p(n);

3 Function Compute-Opt (j):

4 | if j = 0 then

5 | return 0;

6 | else

7 | return

8 | max{v_j+ Compute-Opt(p(j)), Compute-Opt(j - 1)};
```

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Memoization

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

Algorithm 2: Memoization

```
Input: n; s_1, \dots, s_n; f_1, \dots, f_n; v_1, \dots, v_n

1 Sort jobs by finish times so that f_1 \le f_2 \le \dots \le f_n;

2 Compute p(1), p(2), \dots, p(n);

3 for j = 1 \to n do

4 \lfloor M[j] = \text{empty};

5 M[0] = 0;

6 Function M-Compute-Opt (j):

7 \mid \text{if } M[j] \text{ is } empty \text{ then}

8 \mid M[j] = \text{max}\{v_j + \text{M-Compute-Opt}(p(j)), \text{M-Compute-Opt}(j-1)\};

10 \mid \text{return } M[j];
```

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- Sort by finish time: $O(n \log n)$.
- Computing $p(\cdot)$: $O(n \log n)$ via sorting by start time.
- \circ M-Compute-Opt(j): each invocation takes O(1) time and either
 - \triangleright returns an existing value M[j]
 - \triangleright fills in one new entry M[j] and makes two recursive calls
- Progress measure $\Phi =$ number nonempty entries of $M[\cdot]$.
 - \triangleright initially $\Phi = 0$, throughout $\Phi \le n$.
 - \triangleright increases Φ by $1 \Rightarrow$ at most 2n recursive calls.
- Overall running time of M-Compute-Opt(n) is O(n).

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

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Bottom-Up Dynamic Programming

Algorithm 4: Tabulation

Input: n; s_1, \dots, s_n ; f_1, \dots, f_n ; v_1, \dots, v_n

- 1 **Sort** jobs by finish times so that $f_1 \le f_2 \le \cdots \le f_n$;
- **2 Compute** $p(1), p(2), \dots, p(n);$
- 3 Function Iterative-Compute-Opt():
- 4 M[0] = 0;
- for $j = 1 \rightarrow n$ do
- 6 $M[j] = \max\{v_j + M[p(j)], M[j-1]\};$

Finding a Solution from the OPT Value

```
Algorithm 3: Post-Processing

Run M-Compute-Opt(n);

Run Find-Solution(n);

Function Find-Solution (j):

| if j = 0 \text{ then} |

output nothing;

else if v_j + M[p(j)] > M[j-1] \text{ then}

print j;

Find-Solution (p(j));

else

Find-Solution(p(j));
```

∘ # of recursive calls $1 \le n \Rightarrow O(n)$;

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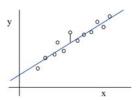
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Four Examples (Recipe of Dynamic Programming)

Segmented Least Squares

- Foundational problem in statistic and numerical analysis.
- Given *n* points in the plane: $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$.
- Find a line y = ax + b to minimize the sum of the squared error:



Solution: Calculus \Rightarrow min error is achieved when

$$a = \frac{n\sum_{i} x_{i} y_{i} - (\sum_{i} x_{i})(\sum_{i} y_{i})}{n\sum_{i} x_{i}^{2} - (\sum_{i} x_{i})^{2}}, b = \frac{\sum_{i} y_{i} - a\sum_{i} x_{i}}{n}$$

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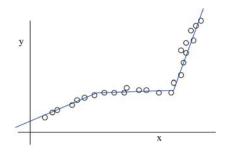
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Segmented Least Squares

- o Points lie roughly on a sequence of several line segments.
- Given *n* points in the plane: $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$ with $x_1 < x_2 < \dots < x_n$, find a sequence of lines that minimizes f(x).

Question: What's a reasonable choice for f(x) to balance accuracy (goodness of fit) and parsimony (number of lines)?



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Multiway Choice

Notation:

- $\circ \ OPT(j) = \text{minimum cost for points } p_1, p_2, \cdots, p_j.$
- \circ $e(i,j) = \text{minimum sum of squares for points } p_i, p_{i+1}, \cdots, p_j.$

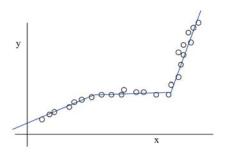
Compute OPT(j):

- Last segment uses points p_i, p_{i+1}, \dots, p_j for some i.
- $\circ \ Cost = e(i,j) + c + OPT(i-1).$

$$OPT(j) = \begin{cases} 0, & j = 0, \\ \min_{1 \le i \le j} \{e(i,j) + c + OPT(i-1)\}, & otherwise \end{cases}$$

Segmented Least Squares

- o Points lie roughly on a sequence of several line segments.
- Given *n* points in the plane: (x_1, y_1) , (x_2, y_2) , \cdots , (x_n, y_n) with $x_1 < x_2 < \cdots < x_n$, find a sequence of lines that minimizes:
 - \triangleright the sum of the sums of the squared errors E in each segment
 - \triangleright the number of lines L
- Tradeoff function: E + cL, for some constant c > 0.



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Segmented Least Squares

Algorithm 5: Tabulation

```
Input: n; p_1, \dots, p_N; c

1 Function Iterative-Compute-Opt():

2 | M[0] = 0;

3 | for j = 1 \rightarrow n do

4 | for i = 1 \rightarrow j do

5 | compute the least square error e_{ij} for the segment

p_i, \dots, p_j;

6 | for j = 1 \rightarrow n do

7 | M[j] = \min_{1 \leq i \leq j} \{e_{ij} + c + M[i - 1]\};

8 | return M[n];
```

Running time: $O(n^3)$ (can be improved to $O(n^2)$ by pre-computing.)

Bottleneck = computing e(i,j) for $O(n^2)$ pairs, O(n) per pair using previous formula.

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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.

Example: $\{3,4\}$ has value 40.

$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$			value	weight
3 18 5		1	1	1
	= 11	2	6	2
4 22 6		3	18	5
		4	22	6
5 28 7		5	28	7

Greedy: repeatedly add item with maximum ratio v_i/w_i .

Example: $\{5, 2, 1\}$ achieves only value = $35 \Rightarrow$ greedy not optimal.

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Adding a New Variable

Definition: $OPT(i, w) = \max \text{ profit subset of items } 1, \dots, i \text{ with weight limit } w.$

Case 1: OPT does not select item *i*.

• OPT selects best of $\{1, 2, \dots, i-1\}$ using weight limit w

Case 2: OPT selects item i.

- new weight $limit = w w_i$
- \circ OPT selects best of using $\{1, 2, \dots, i-1\}$ this new weight limit

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

False Start

Definition: OPT(i)= max profit subset of items $1, \dots, i$.

Case 1: OPT does not select item i.

• OPT selects best of $\{1, 2, \dots, i-1\}$

Case 2: OPT selects item i.

- accepting item *i* does not immediately imply that we will have to reject other items
- without knowing what other items were selected before *i*, we don't even know if we have enough room for *i*

Conclusion: Need more sub-problems!

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Bottom-Up Algorithm (Fill up an *n*-by-*W* array)

Algorithm 6: Knapsack Problem Algorithm using *n*-by-*W* Array

Input:
$$n, W, w_1, \cdots, w_N, v_1, \cdots, v_N$$

1 **for**
$$w = 0 \to W$$
 do
2 | $M[0, w] = 0$;

3 for
$$i = 1 \rightarrow n$$
 do

for
$$w = 1 \rightarrow W$$
 do
if $w_i > w$ then

$$M[i, w] = M[i - 1, w];$$
else

$$M[i, w] = \max\{M[i - 1, w], v_i + M[i - 1, w - w_i]\}$$

9 return M[n, W];

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Knapsack Algorithm

OPT: $\{3, 4\}$ value = 22 + 18 = 40

#	value	weight
1	1	1
2	6	2
3	18	5
4	22	6
5	28	7

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W = 11

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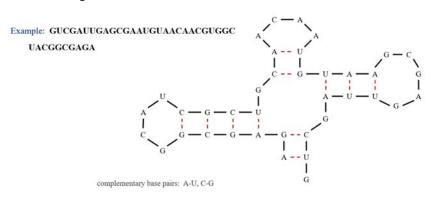
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RNA Secondary Structure

RNA:String $B = b_1 b_2 \cdots b_n$ 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.



Running Time

Running time: $\Theta(nW)$.

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

Knapsack approximation algorithm: There exists a poly-time algorithm that produces a feasible solution that has value within 0.01% of optimum.

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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.

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

Goal: Given an RNA molecule $B = b_1 b_2 \cdots b_n$, find a secondary structure S that maximizes the number of base pairs

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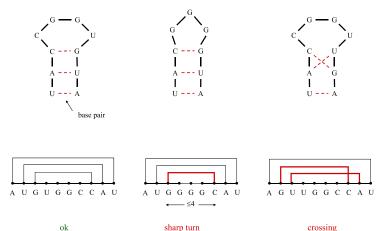
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Examples



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Dynamic Programming Over Intervals

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

Case 1: If $i \ge j - 4$.

• OPT(i,j) = 0 by no-sharp turns condition.

Case 2: Base b_i is not involved in a pair.

 \circ OPT(i,j) = OPT(i,j-1)

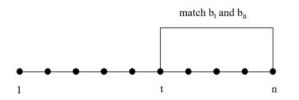
Case 3: Base b_i pairs with b_t for some $i \le t < j - 4$.

- o non-crossing constraint decouples resulting sub-problems
- $\circ OPT(i,j) = 1 + \max_{t} \{ OPT(i,t-1) + OPT(t+1,j-1) \}$

Remark: Same core idea in CKY algorithm to parse context-free grammars.

Subproblems

First attempt: $OPT(j) = \text{maximum number of base pairs in a secondary structure of the substring <math>b_1b_2 \cdots b_j$.



Difficulty: Results in two sub-problems.

- Finding secondary structure in: $b_1b_2 \cdots b_{t-1}$.
- Finding secondary structure in: $b_{t+1}b_{t+2}\cdots b_{n-1}$.

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Bottom Up Dynamic Programming Over Intervals

Question: What order to solve the sub-problems?

Answer: Do shortest intervals first.

Algorithm 7: RNA

Function RNA $(b_1, b_2, \cdots b_n)$:

for $k = 5 \rightarrow n - 1$ do

for $i = 1 \rightarrow n - k$ do j = i + k;
Compute M[i,j]return M[1,n];

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

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String Similarity: How similar are two strings?

Dynamic Programming Summary

Recipe

- o Characterize structure of problem.
- o Recursively define value of optimal solution.
- o Compute value of optimal solution.
- o Construct optimal solution from computed information.

Dynamic programming techniques

- Binary choice: weighted interval scheduling.
- Multi-way choice: segmented least squares.
- Adding a new variable: knapsack.
- o Dynamic programming over interval

Top-down vs. bottom-up: different people have different intuitions.

How similar are two strings? o c c u r r c n c e 6 mismatches, 1 gap o c c u r r a n c e 1 mismatch, 1 gap o c - u r r a n c e

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Basic Methodology String Similarity

Edit Distance

Applications.

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

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

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



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Basic Methodology A Comprehensive Example String Similarity
Sequence Alignment in Linear Space

0 mismatches, 3 gaps

Sequence Alignment

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

Definition: 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.

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

$$M = \sum_{\substack{(x_i, y_j) \in M \\ f \\ \text{mismatch}}} \alpha_{x_i y_j} + \sum_{i: x_i \text{ unmatched}} \delta + \sum_{j: y_j \text{unmatched}} \delta$$

Example: CTACCG vs. TACATG.

Solution: $M = x_2 - y_1, x_3 - y_2, x_4 - y_3, x_5 - y_4,$

 x_6 - y_6 .

Definition: $OPT(i, j) = \min \cos t$ of aligning strings $x_1x_2 \cdots x_i$ and $y_1y_2\cdots y_i$.

Case 1: OPT matches x_i - y_i . pay mismatch for x_i - y_i + min cost of aligning two strings $x_1x_2\cdots x_{i-1}$ and $y_1y_2\cdots y_{i-1}$

Case 2a: OPT leaves x_i unmatched. pay gap for x_i and min cost of aligning $x_1x_2 \cdots x_{i-1}$ and $y_1y_2 \cdots y_i$

Case 2b: OPT leaves y_i unmatched. pay gap for y_i and min cost of aligning $x_1x_2 \cdots x_i$ and $y_1y_2 \cdots y_{i-1}$

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Sequence Alignment

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

English words or sentences: $m, n \le 10$.

Computational biology: m = n = 100,000.10 billions OK, but

10GB array?

Sequence Alignment

Algorithm 8: Sequence Alignment

1 Function

```
Sequence-Alignment (m, n, x_1x_2 \cdots x_m, y_1y_2 \cdots y_n, \delta, \alpha):
        for i = 0 \rightarrow m do
          M[i,0] = i\delta
 3
        for i = 0 \rightarrow n do
          M[0,j] = j\delta
 5
        for i = 1 \rightarrow m do
            for i = 1 \rightarrow n do
 7
                 M[i,j] = \min(\alpha[x_i, y_i] + M[i-1, j-1],
 8
                             \delta + M[i-1,j],
                              \delta + M[i, j-1]
10
        return M[m, n];
11
```

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Sequence Alignment in Linear Space

Linear Space

Question: Can we avoid using quadratic space? Easy. Optimal value in O(m) space and O(mn) time.

- Compute $OPT(i, \cdot)$ from $OPT(i-1, \cdot)$.
- 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.
- Inspired by idea of Savitch from complexity theory.

A Linear Space Algorithm for Computing Maximal Common Subsequences

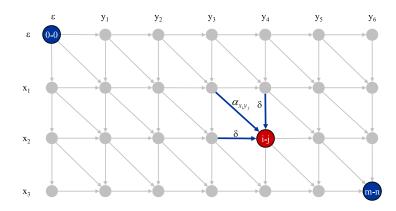
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String Similarity
Sequence Alignment in Linear Space

Edit Distance Graph

• Let f(i,j) be shortest path from (0,0) to (i,j).

• Observation: f(i,j) = OPT(i,j).



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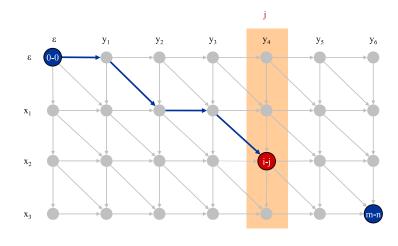
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Sequence Alignment in Linear Space

Edit Distance Graph

- 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) space.



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Edit Distance Graph

 \circ Let f(i,j) be shortest path from (0,0) to (i,j).

• Observation: f(i,j) = OPT(i,j).



Proof: (by strong induction on i + j)

Base case: f(0,0) = OPT(0,0) = 0

Inductive hypothesis: assume true for all (i',j') with i'+j' < i+j.

Induction: Last edge on shortest path to (i,j) is from (i-1,j-1), (i-1,j), or (i,j-1).

$$f(i,j) = \min\{a_{x_iy_i} + f(i-1,j-1), \delta + f(i-1,j), \delta + f(i,j-1)\}$$

= \text{min}\{a_{x_iy_i} + OPT(i-1,j-1), \delta + OPT(i-1,j), \delta + OPT(i,j-1)\}\)
= OPT(i,j)

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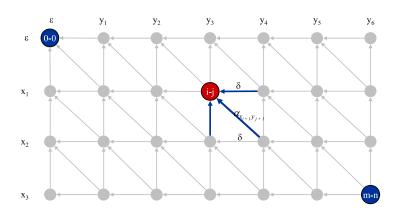
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Sequence Alignment in Linear Space

Edit Distance Graph

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



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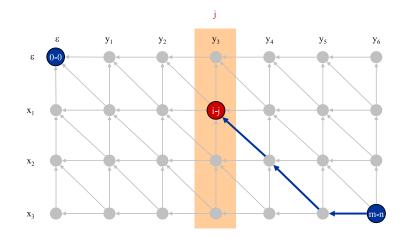
Ovnamic Programming

String Similarity
Sequence Alignment in Linear Space

Edit Distance Graph

• 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) space.



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Dynamic Programming

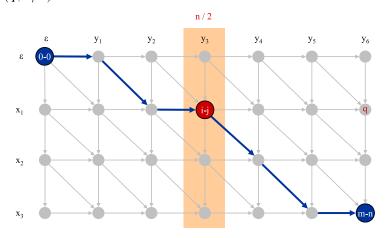
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Basic Methodology A Comprehensive Example

String Similarity
Sequence Alignment in Linear Space

Edit Distance Graph

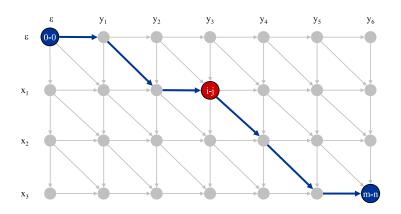
Observation 2: Let q be an index that minimizes f(q, n/2) + g(q, n/2). Then, the shortest path from (0, 0) to (m, n) uses (q, n/2).



Basic Methodology A Comprehensive Example Sequence Alignment in Linear Space

Edit Distance Graph

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



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Basic Methodology A Comprehensive Example

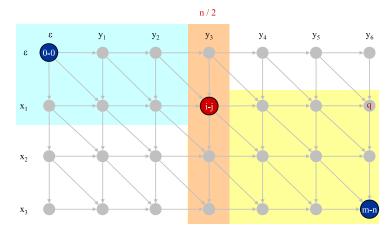
Sequence Alignment in Linear Space

Edit Distance Graph

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

Align x_q and $y_{n/2}$.

Conquer: recursively compute optimal alignment in each piece.



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Running Time Analysis Warmup

Theorem: Let $T(m, n) = \max$ running time of algorithm on strings of length at most m and n. $T(m, n) = O(mn \log n)$.

$$T(m,n) \le 2T(m,n/2) + O(mn) \Rightarrow T(m,n) = O(mn \log n)$$

Remark: Analysis is not tight because two sub-problems are of size (q, n/2) and (m - q, n/2). In next slide, we save $\log n$ factor.

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Basic Methodology A Comprehensive Example

String Similarity
Sequence Alignment in Linear Space

Running Time Analysis (Continued)

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

Proof:

- Base cases: m = 2 or n = 2.
- Inductive hypothesis: $T(m, n) \leq 2cmn$.

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

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

$$= cqn + cmn - cqn + cmn$$

$$= 2cmn$$

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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)

Proof: (by induction on *n*)

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

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

$$T(2,n) \leq cn$$

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

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