Technical Slide

Module 5: Dynamic Programming

Lesson 1: Longest Increasing Subsequence Video 1.1: Warm-up

Video 1.2: Subproblems and Recurrence Relation Video 1.3: Reconstructing a Solution Video 1.4: Subproblems Revisited

2 Lesson 2: Edit Distance

Video 2.1: Algorithm Video 2.2: Reconstructing a Solution Video 2.3: Final Remarks

 Extremely powerful algorithmic technique with applications in optimization, scheduling, planning, economics, bioinformatics, etc

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- Extremely powerful algorithmic technique with applications in optimization, scheduling, planning, economics, bioinformatics, etc
- At contests, probably the most popular type of problems
- A solution is usually not so easy to find, but when found, is easily implementable
- Need a lot of practice!

Fibonacci numbers

Fibonacci numbers

$$F_n = \begin{cases} 0, & n = 0, \\ 1, & n = 1, \\ F_{n-1} + F_{n-2}, & n > 1. \end{cases}$$

 $0, 1, 1, 2, 3, 5, 8, 13, 21, 34, \ldots$

Computing Fibonacci Numbers

Computing F_n

Input: An integer $n \ge 0$. Output: The *n*-th Fibonacci number F_n .

Computing Fibonacci Numbers

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Input: An integer $n \ge 0$. Output: The *n*-th Fibonacci number F_n .

def fib(n):
if
$$n \le 1$$
:
return n
return fib(n - 1) + fib(n - 2)

Recursion Tree



Recursion Tree



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- But Fibonacci numbers grow exponentially fast: $F_n \approx \phi^n$, where $\phi = 1.618...$ is the golden ratio
- E.g., *F*₁₅₀ is already 31 decimal digits long
- The Sun may die before your computer returns
 *F*₁₅₀



Many computations are repeated

Reason

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- "Those who cannot remember the past are condemned to repeat it." (George Santayana)

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- Many computations are repeated
- "Those who cannot remember the past are condemned to repeat it." (George Santayana)
- A simple, but crucial idea: instead of recomputing the intermediate results, let's store them once they are computed

Memoization

```
1 def fib(n):
2 if n <= 1:
3 return n
4 return fib(n - 1) + fib(n - 2)
```

Memoization

```
def fib(n):
1
2
   if n <= 1:
3
        return n
4
    return fib(n - 1) + fib(n - 2)
   T = dict()
1
2
3
   def fib(n):
    if n not in T:
4
5
6
7
8
        if n <= 1:
         T[n] = n
        else:
          T[n] = fib(n - 1) + fib(n - 2)
9
10
     return T[n]
```

But do we really need all this fancy stuff (recursion, memoization, dictionaries) to solve this simple problem?

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$$F_0 = 0, F_1 = 1$$

2 $F_2 = 0 + 1 = 1$
3 $F_3 = 1 + 1 = 2$
4 $F_4 = 1 + 2 = 3$

- But do we really need all this fancy stuff (recursion, memoization, dictionaries) to solve this simple problem?
- After all, this is how you would compute F₅ by hand:

1
$$F_0 = 0, F_1 = 1$$

2 $F_2 = 0 + 1 = 1$
3 $F_3 = 1 + 1 = 2$
4 $F_4 = 1 + 2 = 3$
5 $F_5 = 2 + 3 = 5$

Iterative Algorithm

def fib(n):

$$T = [None] * (n + 1)$$

$$T[0], T[1] = 0, 1$$
for i in range(2, n + 1):

$$T[i] = T[i - 1] + T[i - 2]$$
return T[n]



But do we really need to waste so much space?

Hm Again...

But do we really need to waste so much space?

```
def fib(n):
1
   if n <= 1:
2
3
4
5
6
7
       return n
     previous, current = 0, 1
     for in range(n - 1):
       new current = previous + current
8
       previous, current = current, new current
9
10
     return current
```

• O(n) additions

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- On the other hand, recall that Fibonacci numbers grow exponentially fast: the binary length of F_n is O(n)

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- On the other hand, recall that Fibonacci numbers grow exponentially fast: the binary length of F_n is O(n)
- In theory: we should not treat such additions as basic operations
- In practice: just F₁₀₀ does not fit into a 64-bit integer type anymore, hence we need bignum arithmetic



The key idea of dynamic programming: avoid recomputing the same thing again!

Summary

- The key idea of dynamic programming: avoid recomputing the same thing again!
- At the same time, the case of Fibonacci numbers is a slightly artificial example of dynamic programming since it is clear from the very beginning what intermediate results we need to compute the final result
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Longest Increasing Subsequence

Longest increasing subsequence

Input: An array $A = [a_0, a_1, \dots, a_{n-1}]$. Output: A longest increasing subsequence (LIS), i.e., $a_{i_1}, a_{i_2}, \dots, a_{i_k}$ such that $i_1 < i_2 < \dots < i_k, a_{i_1} < a_{i_2} < \dots < a_{i_k}$, and k is maximal.









 Consider the last element x of an optimal increasing subsequence and its previous element z:



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- Moreover, the prefix of the IS ending at z must be an optimal IS ending at z as otherwise the initial IS would not be optimal:



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Optimal substructure by "cut-and-paste" trick

 Let LIS(i) be the optimal length of a LIS ending at A[i]

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- Then

 $LIS(i) = 1 + \max\{LIS(j) : j < i \text{ and } A[j] < A[i]\}$

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 Convention: maximum of an empty set is equal to zero

 Let LIS(i) be the optimal length of a LIS ending at A[i]

Then

 $LIS(i) = 1 + \max\{LIS(j) : j < i \text{ and } A[j] < A[i]\}$

- Convention: maximum of an empty set is equal to zero
- Base case: *LIS*(0) = 1

Algorithm

When we have a recurrence relation at hand, converting it to a recursive algorithm with memoization is just a technicality

We will use a table T to store the results:
 T[i] = LIS(i)

Algorithm

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- We will use a table T to store the results: T[i] = LIS(i)
- Initially, T is empty. When LIS(i) is computed, we store its value at T[i] (so that we will never recompute LIS(i) again)

Algorithm

When we have a recurrence relation at hand, converting it to a recursive algorithm with memoization is just a technicality

- We will use a table T to store the results: T[i] = LIS(i)
- Initially, T is empty. When LIS(i) is computed, we store its value at T[i] (so that we will never recompute LIS(i) again)
- The exact data structure behind T is not that important at this point: it could be an array or a hash table

Memoization

```
\top = dict()
1
2
3
   def lis (A, i):
4
    if i not in T:
5
6
7
8
       T[i] = 1
        for j in range(i):
          if A[j] < A[i]:
9
            T[i] = \max(T[i], \text{ lis}(A, i) + 1)
10
     return T[i]
11
12
13
   A = [7, 2, 1, 3, 8, 4, 9, 1, 2, 6, 5, 9, 3]
   print(max(lis(A, i) for i in range(len(A))))
14
```

Running Time

The running time is quadratic $(O(n^2))$: there are *n* "serious" recursive calls (that are not just table look-ups), each of them needs time O(n) (not counting the inner recursive calls)

Table and Recursion

■ We need to store in the table T the value of LIS(i) for all i from 0 to n - 1

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Table and Recursion

- We need to store in the table T the value of LIS(i) for all i from 0 to n - 1
- Reasonable choice of a data structure for *T*: an array of size *n*
- Moreover, one can fill in this array iteratively instead of recursively

Iterative Algorithm



Iterative Algorithm



 Crucial property: when computing T[i], T[j] for all j < i have already been computed

Iterative Algorithm



Crucial property: when computing T[i], T[j] for all j < i have already been computed
 Durning time: Q(n²)

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Reconstructing a Solution

■ How to reconstruct an optimal IS?

Reconstructing a Solution

- How to reconstruct an optimal IS?
- In order to reconstruct it, for each subproblem we will keep its optimal value and a choice leading to this value

Adjusting the Algorithm

```
def lis(A):
1
2
3
4
5
6
7
8
     T = [None] * Ien(A)
      prev = [None] * len(A)
      for i in range(len(A)):
        T[i] = 1
        prev[i] = -1
        for j in range(i):
9
           if A[j] < A[i] and T[i] < T[j] + 1:
             T[i] = T[i] + 1
10
             prev[i] = i
11
```






























































Example



Unwinding Solution

```
1
      last = 0
2
3
4
5
6
7
      for i in range(1, \text{len}(A)):
        if T[i] > T[last]:
           last = i
      lis = []
      current = last
8
9
      while current >= 0:
        lis.append(current)
        current = prev[current]
10
      lis.reverse()
11
      return [A[i] for i in lis]
12
```













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- A recurrence relation for subproblems can be immediately converted into a recursive algorithm with memoization
- A recursive algorithm, in turn, can be converted into an iterative one
- An optimal solution can be recovered either by using an additional bookkeeping info or by using the computed solutions to all subproblems

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- In most DP algorithms, the most creative part is coming up with the right notion of a subproblem and a recurrence relation
- When a recurrence relation is written down, it can be wrapped with memoization to get a recursive algorithm
- In the previous video, we arrived at a reasonable subproblem by analyzing the structure of an optimal solution
- In this video, we'll provide an alternative way of arriving at subproblems: implement a naive brute force solution, then optimize it

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Start with an empty sequence

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 - Start with an empty sequence
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- Need the longest increasing subsequence? No problem! Just iterate over all subsequences and select the longest one:
 - Start with an empty sequence
 - Extend it element by element recursively
 - Keep track of the length of the sequence
- This is going to be slow, but not to worry: we will optimize it later

Brute Force: Code

```
def lis (A, seq):
1
     result = len(seq)
2
3
4
5
6
     if len(seq) == 0:
       last index = -1
        last element = float("-inf")
7
     else:
8
        last index = seq[-1]
9
        last element = A[last index]
10
11
     for i in range(last index + 1, len(A)):
12
        if A[i] > last element:
13
          result = max(result, lis(A, seq + [i]))
14
     return result
15
16
17
   print(lis(A=[7, 2, 1, 3, 8, 4, 9], seq=[]))
```



 At each step, we are trying to extend the current sequence

Optimizing

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Optimizing

- At each step, we are trying to extend the current sequence
- For this, we pass the current sequence to each recursive call
- At the same time, code inspection reveals that we are not using all of the sequence: we are only interested in its last element and its length
- Let's optimize!

Optimized Code

1 2

3

4 5

6 7 8

9

10

11

12 13

14 15

16

```
def lis (A, seq len, last index):
  if last index = -1:
    last element = float("-inf")
  else:
    last element = A[last index]
  result = seq len
  for i in range(last index + 1, len(A)):
    if A[i] > last element:
      result = max(result,
                lis(A, seq len + 1, i))
  return result
print(lis([3, 2, 7, 8, 9, 5, 8], 0, -1))
```

Inspecting the code further, we realize that seq_len is not used for extending the current sequence (we don't need to know even the length of the initial part of the sequence to optimally extend it)

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- More formally, for any x, extend(A, seq_len, i) is equal to extend(A, seq_len - x, i) + x

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- More formally, for any x, extend(A, seq_len, i) is equal to extend(A, seq_len - x, i) + x
- Hence, can optimize the code as follows: max(result, 1 + seq_len + extend(A, 0, i))

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- More formally, for any x, extend(A, seq_len, i) is equal to extend(A, seq_len - x, i) + x
- Hence, can optimize the code as follows: max(result, 1 + seq_len + extend(A, 0, i))
- Excludes seq_len from the list of parameters!

Resulting Code

1

10

11 12

```
def lis (A, last index):
2
    if last index == -1:
3
       last element = float("-inf")
4
     else:
5
       last element = A[last index]
6
7
8
9
     result = 0
     for i in range(last index + 1, len(A)):
       if A[i] > last element:
          result = max(result, 1 + lis(A, i))
13
     return result
14
   print(lis([8, 2, 3, 4, 5, 6, 7], -1))
15
```

Resulting Code

```
def lis (A, last index):
1
     if last index == -1:
2
3
        last element = float("-inf")
4
5
     else:
        last element = A[last index]
6
7
8
9
     result = 0
     for i in range(last index + 1, len(A)):
        if A[i] > last element:
10
          result = max(result, 1 + lis(A, i))
11
12
13
     return result
14
   print(lis([8, 2, 3, 4, 5, 6, 7], -1))
15
   It remains to add memoization!
```


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- Two common ways of arriving at the right subproblem:



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- Two common ways of arriving at the right subproblem:
 - Analyze the structure of an optimal solution

Summary

- Subproblems (and recurrence relation on them) is the most important ingredient of a dynamic programming algorithm
- Two common ways of arriving at the right subproblem:
 - Analyze the structure of an optimal solutionImplement a brute force solution and optimize it

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Statement

Edit distance

Input: Two strings $A[0 \dots n-1]$ and $B[0 \dots m-1]$.

Output: The minimal number of insertions, deletions, and substitutions needed to transform A to B. This number is known as edit distance or Levenshtein distance.

EDITING

EDITING ↓remove E DITING

EDITING remove E DITING insert S DISTING replace I with by A DISTANG replace G with C DISTANC

EDITING remove E DITING insert S DISTING replace I with by A DISTANG replace G with C DISTANC insert E DISTANCE

Example: alignment



cost: 5

Example: alignment

substitutions/mismatches



$$A[0\ldots n-1]$$
$$B[0\ldots m-1]$$







Subproblems

• Let ED(i, j) be the edit distance of $A[0 \dots i - 1]$ and $B[0 \dots j - 1]$.

Subproblems

- Let ED(i, j) be the edit distance of $A[0 \dots i 1]$ and $B[0 \dots j 1]$.
- We know for sure that the last column of an optimal alignment is either an insertion, a deletion, or a match/mismatch.

Subproblems

- Let ED(i, j) be the edit distance of A[0...i - 1] and B[0...j - 1].
- We know for sure that the last column of an optimal alignment is either an insertion, a deletion, or a match/mismatch.
- What is left is an optimal alignment of the corresponding two prefixes (by cut-and-paste).

Recurrence Relation

$$ED(i, j) = \min \begin{cases} ED(i, j - 1) + 1\\ ED(i - 1, j) + 1\\ ED(i - 1, j - 1) + diff(A[i], B[j]) \end{cases}$$

Recurrence Relation

$$ED(i, j) = \min \begin{cases} ED(i, j - 1) + 1 \\ ED(i - 1, j) + 1 \\ ED(i - 1, j - 1) + diff(A[i], B[j]) \end{cases}$$

Base case: ED(i, 0) = i, ED(0, j) = j

Recursive Algorithm

```
T = dict()
def edit distance(a, b, i, j):
  if not (i, j) in T:
    if i == 0: T[i, j] = j
    elif i == 0: T[i, j] = i
    else:
      diff = 0 if a[i - 1] == b[j - 1] else 1
      T[i, j] = min(
        edit distance (a, b, i - 1, j) + 1,
        edit distance (a, b, i, j - 1) + 1,
        edit distance(a, b, i - 1, j - 1) + diff)
  return T[i, j]
print(edit distance(a="editing", b="distance",
                     i = 7, i = 8)
```

Converting to a Recursive Algorithm

Use a 2D table to store the intermediate results

Converting to a Recursive Algorithm

- Use a 2D table to store the intermediate results
- ED(i,j) depends on ED(i-1,j-1), ED(i-1,j), and ED(i,j-1):



Filling the Table

Fill in the table row by row or column by column:





Iterative Algorithm

```
def edit distance(a, b):
1
2
     T = [[float("inf")] * (len(b) + 1)]
3
           for in range(len(a) + 1)]
4
     for i in range(len(a) + 1):
5
      T[i][0] = i
6
7
8
     for j in range(len(b) + 1):
       T[0][i] = i
9
     for i in range (1, \text{len}(a) + 1):
10
        for j in range(1, len(b) + 1):
          diff = 0 if a[i - 1] == b[j - 1] else 1
11
         T[i][j] = min(T[i - 1][j] + 1,
12
13
                        T[i][i - 1] + 1,
                        T[i - 1][i - 1] + diff
14
15
16
     return T[len(a)][len(b)]
17
18
   print(edit distance(a="distance", b="editing"))
19
```
















Brute Force

Recursively construct an alignment column by column

Brute Force

- Recursively construct an alignment column by column
- Then note, that for extending the partially constructed alignment optimally, one only needs to know the already used length of prefix of A and the length of prefix of B

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- If ED(i,j) = ED(i 1, j) + 1, then there exists an optimal alignment whose last column is a deletion

- To reconstruct a solution, we go back from the cell (n, m) to the cell (0, 0)
- If ED(i,j) = ED(i 1, j) + 1, then there exists an optimal alignment whose last column is a deletion
- If ED(i, j) = ED(i, j 1) + 1, then there exists an optimal alignment whose last column is an insertion

- To reconstruct a solution, we go back from the cell (n, m) to the cell (0, 0)
- If ED(i, j) = ED(i 1, j) + 1, then there exists an optimal alignment whose last column is a deletion
- If ED(i,j) = ED(i,j-1) + 1, then there exists an optimal alignment whose last column is an insertion
- If ED(i,j) = ED(i-1,j-1) + diff(A[i], B[j]), then match (if A[i] = B[j]) or mismatch (if $A[i] \neq B[j]$)







deletion



С	Ε
-	G

deletion



Ν	С	Ε
Ν	-	G























Technical Slide

Module 5: Dynamic Programming

Lesson 1: Longest Increasing Subsequence Video 1.1: Warm-up Video 1.2: Subproblems and Recurrence Relati Video 1.3: Reconstructing a Solution Video 1.4: Subproblems Revisited

2 Lesson 2: Edit Distance

Video 2.1: Algorithm Video 2.2: Reconstructing a Solution Video 2.3: Final Remarks

Saving Space

When filling in the matrix it is enough to keep only the current column and the previous column:



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Thus, one can compute the edit distance of two given strings A[1...n] and B[1...m] in time O(nm) and space O(min{n, m}).

 However we need the whole table to find an actual alignment (we trace an alignment from the bottom right corner to the top left corner)

- However we need the whole table to find an actual alignment (we trace an alignment from the bottom right corner to the top left corner)
- There exists an algorithm constructing an optimal alignment in time O(nm) and space O(n + m) (Hirschberg's algorithm)

Weighted Edit Distance

- The cost of insertions, deletions, and substitutions is not necessarily identical
- Spell checking: some substitutions are more likely than others
- Biology: some mutations are more likely than others

Generalized Recurrence Relation

$$\min \begin{cases} ED(i, j - 1) + \text{inscost}(B[j]), \\ ED(i - 1, j) + \text{delcost}(A[i]), \\ ED(i - 1, j - 1) + \text{substcost}(A[i], B[j]) \end{cases}$$