Week 04: Alignment

 ${\sf FanFly}$

March 29, 2020

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- The distance is calculated by counting the minimum number of operations needed to transform one string into the other.
- Different definitions of an edit distance use different sets of string operations.

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- We can transform weather into whether by inserting an h and deleting an a, and thus the edit distance between them is 2.
- Note that if we remove the different parts from both strings, we will get their longest common subsequence, i.e., wether.
- Conversely, if we can find their longest common subsequence, we can calculate their edit distance.



Now we introduce the longest common subsequence problem.

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The Longest Common Subsequence Problem

- Input: Two sequences A and B of lengths n and m, respectively.
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mea--surement -measur-ement --amus--ement am---u-sement
```

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• If x = y, then we have

$$LCS(A, B) = LCS(A', B') \cdot x.$$

Otherwise, we have

$$LCS(A, B) = LCS(A, B')$$
 or $LCS(A, B) = LCS(A', B)$.

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Recursive Method

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```
def lcs(A, B, n, m):
if n == 0 or m == 0:
   return A[:0]
elif A[n - 1] == B[m - 1]:
    return lcs(A, B, n - 1, m - 1) + A[n - 1]
else:
    P = lcs(A, B, n - 1, m)
    Q = lcs(A, B, n, m - 1)
    if len(P) > len(Q):
       return P
    else:
        return 0
```

Time Complexity of Recursive Method

What is its time complexity?



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$$T(n,m) = \begin{cases} O(1), & \text{if } n = 0 \text{ or } m = 0 \\ T(n,m-1) + T(n-1,m) + O(1), & \text{otherwise.} \end{cases}$$



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Thus, we can conclude that

$$T(n,m) = \Theta\left(\frac{(n+m)!}{n! \cdot m!}\right),$$

which is really inefficient.



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- If the algorithm can remember the solutions to the solved subproblems, then it can become more efficient.

	m	a	s	k
i	-	-	-	_
d	_	-	-	-
е	-	-	-	-
a	_	а	a	a
s	-	a	as	as

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S	-	a	as	as

• This improvement is called **dynamic programming**.

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- Assume that the length of longest common subsequence is ℓ .
- The content of each entry can be computed in $\Theta(\ell)$ time.
- Thus, we can solve the problem in $O(nm\ell)$ time.



Improvement

In fact, the time complexity can be improved to O(nm) if we only memoize the length of the longest common subsequence.

	m	a	s	k
i	0	0	0	0
d	0	0	0	0
е	0	0	0	0
a	0	1	1	1
s	0	1	2	2

Let A and B be sequences of lengths n and m, respectively.

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Solution

The edit distance between A and B is $(n - \ell) + (m - \ell) = n + m - 2\ell$.



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Solution

It has $2^n - 1$ nonempty subsequences.



Let A and B be sequences and let f be a scoring function. Suppose that

$$f(x,y) = \begin{cases} p, & \text{if } x = y \\ q, & \text{otherwise.} \end{cases}$$



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$$f(x,y) = \begin{cases} p, & \text{if } x = y \\ q, & \text{otherwise.} \end{cases}$$

For any alignment, we can compute its score according to the scoring function.



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Exercise #3 (cont.)

For example, the alignment below has score 3p + 6q.

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Please determine the value of p and q such that the maximum score of alignment of A and B is equal to the length of their longest common subsequence.



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For example, the alignment below has score 3p + 6q.

Please determine the value of p and q such that the maximum score of alignment of A and B is equal to the length of their longest common subsequence.

Solution

This property is satisfied when p=1 and q=0.



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