



Longest Common Subsequences

Seminar 2

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May 31, 2021

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1. Introduction

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1.1 What are LCS ?

What are LCS ?

Notation

“LCS” = Longest Common Subsequence(s)

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Example 1

S_1 : A B A B B

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S_1 : A B A B B

S_2 : A A B A B

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S_1 : **A** B A B B

S_2 : **A** A B A B

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S_1 : **A** **B** A B B
 S_2 : **A** A **B** A B

What are LCS ?

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Example 1

S_1 :	A	B	A	B	B
S_2 :	A	A	B	A	B

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S_1 : **A** **B** **A** **B** B

S_2 : **A** A **B** **A** **B**

\Rightarrow The LCS between S_1 and S_2 is **A B A B**

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Notation

“LCS” = Longest Common Subsequence(s)

Example 1

S_1 : **A** **B** **A** **B** B
 S_2 : **A** A **B** **A** **B**

\Rightarrow The LCS between S_1 and S_2 is **A B A B**

*NB: LCS may not be unique, **A A B B** also works.*

Example 2

What is the LCS of the following sequences ?

Example 2

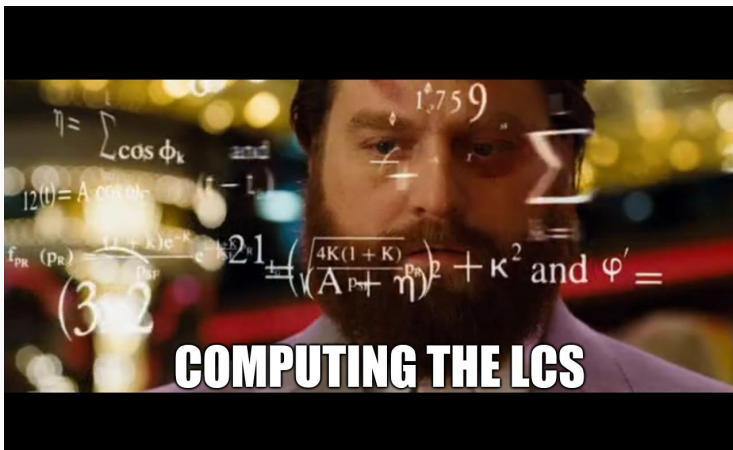
What is the LCS of the following sequences ?

S_3 : AABBAABAAABABAAABAAABBABAAABAAABAAA
AAAABBABBBBAAABABBAABBAABBBBBBAAAABA
BBABAAAABABAABBBBABBBBBBAAABBBBAABBB
AABAABBABABAABABBBBBBBAABBBBBBAAAAAB
AABAAAAABAABAABAAABBABBBBABBAABBB

S_4 : BABBBABAABAABBBABBABBBBBBBBABABAAABB
BBABBABBABBBABBBABBABBABABABBAABABA
BAABABAAAABABBABABBAABABBAABABABB
BABBBBBBBAABAABBBABBBBBAAAAABBBBBBAAAB
ABBAAAABBBBABABAABBABBBBAABABBBABAABA

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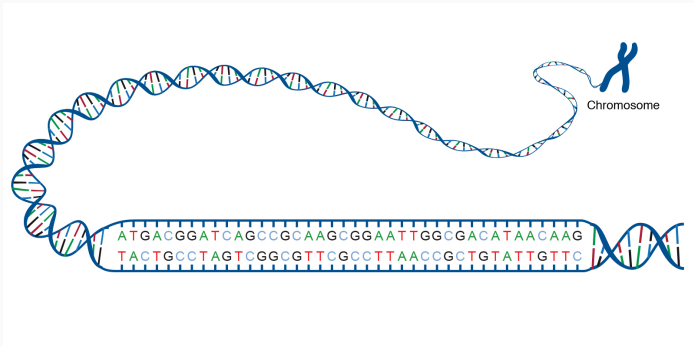
1. Introduction

1.2 Why are we interested in LCS ?

Applications

Applications:

- Bioinformatics: Compare sequences of nucleotides (DNA)



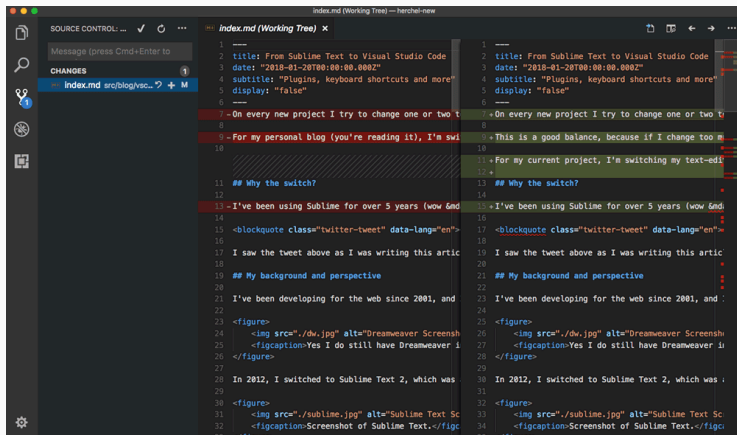
Applications:

- Bioinformatics: Compare sequences of nucleotides (DNA)
- Natural Language Processing: Compare texts



Applications:

- Bioinformatics: Compare sequences of nucleotides (DNA)
- Natural Language Processing: Compare texts
- Computer Science: Spot differences in texts



2. How to find LCS ?

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2.1 Step 1: Building the table

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2.2 Step 2: Crawling back up the table

3. Data analysis of LCS results

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3.1 Subsection 1

Thank you