

# On Solving the Multiple Variable Gapped Longest Common Subsequence Problem

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# Outline

- Introduction & Preliminaries
- Problem Definition
- Graph state space
- Iterative multi-source Beam Search
- Experimental Evaluation
  - General problem
  - Special problem
- Conclusions

# Introduction

- Objects we deal with: sequences (strings) over finite alphabet
  - DNA/RNA over {A, T, G, C/U}
  - Proteins over 20 (canonical) amino acids: {A, C, D, E, P, Q...}
- Computational biology
  - **One of central tasks:** sequence comparison, finding common motifs between sequences
  - compare structurally but also semantically/functionality
  - sequence alignment problems
- Subsequences: reveal structural similarities → **Longest common subsequence problem** variants

# Longest common subsequence problem (LCSP)

- Basic problem in Computational biology
- Intensively solved over last 50 years
  - theoretically as well as practically
  - Practically: many approximation algorithms, (meta-) heuristics, exact approaches, etc.

## Definition (LCSP)

**Input:** Given a set of sequences  $S = \{s_1, \dots, s_m\}$

**Task:** Find a subsequence  $s$  which is **common** for all sequences from  $S$  of **maximum** possible length.

## Example

Input:  $S = \{\text{AATTGC}, \text{ATTAC}\}$

LCS solution:  $s = \text{ATTC}$

# Literature & LCS Problem Variants

Theoretically:

- When  $m = 2$  – polynomially solvable (in  $O(n^2)$ ): Dynamic programming, Hunt-Schlimanski, ...
- When  $m$  arbitrary large –  $\mathcal{NP}$ -hard:
  - subject of interest within last 30 years: approximation approaches, meta-heuristics (ACO, Beam search, ...), but also exact approaches (A\*, anytime approaches,...)

Problem-related variants:

- Arc-annotated LCS problem
- Constrained (restricted/imposed) LCS problem
- Repetition-free, Longest filled LCS,...
- **Gapped LCS problem**

# The gaped LCS problem

## Definition (A gap sequence)

Given is a sequence  $s$  and an assigned function  $G_s: \{1, \dots, |s|\} \mapsto \mathbb{N}$ . An ordered pair  $(s, G_s)$  is called a **sequence with gaps**.

## Definition (A gapped subsequence )

Sequence  $\tilde{s}$  is a gapped subsequence of  $(s, G_s)$  iff

- $\tilde{s}$  is a subsequence of  $s$
- the gapped constraint  $G_s$  is fulfilled

# Literature & Motivation

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