

# Advanced Algorithms in Bioinformatics (P4)

## Sequence and Structure Analysis

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### *Exercise 1.*

#### Efficient searching with suffix arrays

In the lecture we discussed two strategies how to reduce the number of redundant character comparisons during a binary search. One uses the mlr values, while the other one makes use of lcp values. The mlr trick in practice already brings the running time to  $O(m + \log n)$ .

- Find a pair of pattern and text where the mlr trick still needs time  $O(m \log n)$ .
- For the same text and pattern perform the binary search using the lcp values.
- Prove that using the lcp method the search algorithm does at most  $O(m + \log n)$  character comparisons.

### *Exercise 2.*

Given a text  $T$  of length  $n$ , let  $suftab'$  be the suffix array of  $T$  where suffixes are lexicographically ordered according to the first  $m$  letters for some  $m < n$ . Will the Kasai algorithm still compute the correct lcp values of adjacent suffixes in  $suftab'$ ? Justify your answer!

### *Exercise 3.*

Given the text **halloballo** construct the suffix array using the Skew algorithm.

### *Exercise 4.*

Show that the worst case runtime of the skew algorithm for a text of length  $n$  is  $O(n)$ .