

# Homework 5

CS 4364/5364  
Spring 2021

Due: 29 April 2021

1. Given an additive tree  $T = (E, V)$  for  $n$  species. (a) Describe an algorithm for reconstructing the distance matrix between all of the species. (b) What is the time complexity of the algorithm you described in (a) algorithm?
2. Given a set of sequences  $R = (r_1, r_2, r_3, \dots, r_\ell)$ , which may contain single character changes (i.e. it may be one character from the alphabet, but should have been another). Design an algorithm that outputs a modified set of reads  $R' = (r'_1, r'_2, r'_3, \dots, r'_\ell)$  that replaces any changes such that the number of  $k$ -mers in  $R'$  that are erroneous is lowered (it may not be eliminated, we define this below).

An *erroneous*  $k$ -mer is one that occurs at least once and less than 5 times. Note that one character change will impact up to  $2k - 1$  overlapping  $k$ -mers.

You can assume you have access to a  $k$ -mer conversion function  $f(x) = y$  such that assigns an integer  $y \in [1 \dots \sigma^k]$  to each  $x \in \Sigma^k$ , and a  $k$ -mer count array  $C[0 \dots \sigma^k]$  where  $C[y]$  contains the number of times  $f^{-1}(y)$  occurs in  $R$ . (Here  $f^{-1}(y)$  returns the  $k$ -mer  $x$  given an index  $y$ .)

You can also assume that any window of  $2k$  bases will only have 1 error, i.e. there will never be conflicts where two point mutations in the same  $k$ -mer. In the case that a character could be replaced with two different characters and satisfy this condition, prefer the one that has more total occurrences across the  $k$  overlapping windows.

**Note:** There are multiple solutions, some examples include a greedy solution, a dynamic programming solution, and an ILP. You can choose any of these as long as you justify that your solution will not *increase* the number of erroneous  $k$ -mers.

**Example:** assume  $k = 3$  and the following read segment corrections would be made in this window of 6 characters given these  $k$ -mer frequencies:

...ACTTG...  $\longrightarrow$  ...ACCTG...

$x$	$C[f(x)]$
ACA	100
ACC	50 $\rightarrow$ 51
ACT	1 $\rightarrow$ 0
ACG	9
ATG	2
CAT	4
CCT	12 $\rightarrow$ 13
CTT	4 $\rightarrow$ 3
CTG	7 $\rightarrow$ 8
CGT	0
TTG	2 $\rightarrow$ 1
GTG	3

In this example, changing the middle T to a C eliminates three occurrences of erroneous  $k$ -mers. The change in any values in the count array are illustrated, the original value is to the left, the counts after the change are shown to the right.