

# Bioinformatics Algorithms

Lecture Notes

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# 1 Definitions

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## 1.1 Alphabet

Let  $\Sigma$  be a finite set of symbols (alternatively called characters), called the alphabet. No assumption is made about the nature of the symbols.

## 1.2 Strings

A string (or word) over  $\Sigma$  is any finite sequence of symbols from  $\Sigma$ . For example if  $\Sigma = \{0, 1\}$  then 01011 is a string over  $\Sigma$ .

The **length of a string  $S$  is the number of symbols in  $s$**  (the length of the sequence) and can be any non-negative integer. It is often denoted as  $|S|$ .

The **empty string** is the unique string over  $\Sigma$  of length 0, and is denoted as  $\epsilon$ .

The **set of all strings over  $\Sigma$  of length  $n$**  is denoted  $\Sigma^n$ . For example, if  $\Sigma = 0, 1$ , then  $\Sigma^2 = \{00, 01, 10, 11\}$ . Note that  $\Sigma^0 = \{\epsilon\}$  for any alphabet  $\Sigma$ .

The **set of all strings of any length over  $\Sigma$**  is the *Kleene closure* of  $\Sigma$  and is denoted as  $\Sigma^*$ . Also:

$$\Sigma^* = \bigcup_{n \in \mathbb{N} \cup \{0\}} \Sigma^n$$

The **set of all non-empty strings over  $\Sigma$**  is denoted by  $\Sigma^+$

## 1.3 Subsequences

A **subsequence** of  $S$  is a string that can be **obtained by deleting any number (from 0 to  $n$ ) of non-consecutive characters** from  $S$ , including  $S$  and the empty string too.

### 1.3.1 Let us count: subsequences

A subsequence of  $S$  can be described by a binary string  $B$  of  $n$  elements, telling us for each position  $i$  if the corresponding character is kept in the subsequence (1) or deleted (0).

- $S = \text{apple}, B = 11010 \rightarrow \text{subsequence} = \text{apl}$

Thus, the **number of subsequences** corresponds to the number of binary strings of length  $n$ , hence

$$\text{subsequences} = 2^n$$

Notice that if two subsequences produced by different choices are **identical, they will be counted twice**.

## 1.4 Substrings

A string  $s$  is said to be a **substring** (or *factor*) of  $t$  if there exist (possibly empty) strings  $u$  and  $v$  such that  $t = usv$ .

Given a string  $t$ , **suffixes** and **prefixes** are special substrings of  $t$ .

### 1.4.1 Let us count: substrings

Given a string of length  $n$ :

- We have 1 substring of length  $n$  (starting at 1)
- We have 2 substrings of length  $n-1$  (starting at 1 and 2)
- We have 3 substrings of length  $n-2$  (starting at 1,2 and 3)
- ...
- We have  $n - i + 1$  substrings of length  $i$  (starting at all positions from 1 to  $n - i + 1$ )
- We have  $n$  substrings of length 1 (starting at every position)

Plus, 1 substring of length 0.

This brings the **total number of substrings** of a string of length  $n$  to

$$\text{substrings} = \sum_{i=1}^n i = \frac{n(n+1)}{2} + 1$$

## 1.5 Prefixes and suffixes

A string  $s$  is said to be a **prefix** of  $t$  if there exists a string  $u$  such that  $t = su$ . If  $u$  is nonempty,  $s$  is said to be a *proper* prefix of  $t$ .

Simmetrically, a string  $s$  is said to be a **suffix** of  $t$  if there exists a string  $u$  such that  $t = us$ . If  $u$  is nonempty,  $s$  is said to be a *proper* suffix of  $t$ .

### 1.5.1 Let us count: prefixes and suffixes

The number of all possible prefixes/suffixes of a string of length  $n$  is

$$\text{prefixes/suffixes} = n + 1$$

$n$  are non-empty,  $+1$  because we include also the **empty string** as prefix/suffix.

## 1.6 Reverse and palindrome

The **reverse** of a string is a string with the same symbols but in reverse order. For example, if  $s = abc$  (where  $a$ ,  $b$ , and  $c$  are symbols of the alphabet), then the reverse of  $s$  is  $cba$ .

A string that is the reverse of itself is called a **palindrome**, which also includes the empty string and all strings of length 1.

## 1.7 Rotations and permutations

A string  $s = uv$  is said to be a **rotation** of  $t$  if  $t = vu$ . For example, if  $\Sigma = \{0,1\}$  the string  $0011001$  is a rotation of  $0100110$ , where  $u = 00110$  and  $v = 01$ .

### 1.7.1 Let us count: rotations

Given a string  $S$ , with  $|S| = n$ , we denote  $S=AB$ , where **A is a prefix** substring, and **B a suffix** substring.

For every possible pair  $A, B$  the string  **$R=BA$  is a rotation of  $S$** , including  $S$  itself, that is, the case where  $A = \epsilon$  and  $B = S$ .

Since we have  $n$  different non empty suffixes  $B$ , the number of rotations is

$$rotations = n$$

It is not  $n + 1$  because the two suffixes  $B = S$  and  $B = \epsilon$  produce exactly the same rotation, given by  $S$  itself.

### 1.7.2 Let us count: permutations

Given a string  $S$ , with  $|S| = n$ , the number of permutations of the characters in  $S$  is

$$permutations = n!$$

Notice that the formula does not consider the case of identical permutations: that is, a string of  $n$  identical characters, will have  $n$  identical permutations.

## 2 Comparing strings

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### 2.1 Hamming distance

The **Hamming distance** between **two strings of equal length** is the **number of positions at which the corresponding symbols are different**.

- “karolin” and “kathrin”  $\rightarrow 3$
- 1011101 and 1001001  $\rightarrow 2$

With Hamming distance we can formalize *substitutions* in biological sequences - or simply sequencing errors in which the wrong base pair is identified.

### 2.2 Edit distance

The **edit distance** is a way of quantifying how dissimilar two strings (e.g., words) are to one another by counting the **minimum number of operations required to transform one string into the other**. In the *Levenshtein distance* (the most common), edit operations are: **removal**, **insertion**, and **substitution**.

The edit distance between “kitten” and “sitting” is 3. A minimal edit script that transforms the former into the latter is:

- kitten  $\rightarrow$  sitten (substitute *s* for *k*)
- sitten  $\rightarrow$  sittin (substitute *i* for *e*)
- sittin  $\rightarrow$  sitting (insert *g* at the end)

The number of solutions (sequences of operations) is infinite.