### **Bioinformatics Algorithms**

(Fundamental Algorithms, module 2)

#### Zsuzsanna Lipták

Masters in Medical Bioinformatics academic year 2018/19, II semester

Strings and Sequences in Computer Science

• ∑ a finite set called alphabet

- ∑ a finite set called alphabet
- its elements are called characters or letters

- Σ a finite set called alphabet
- its elements are called characters or letters
- $|\Sigma|$  is the size of the alphabet (number of different characters)

- Σ a finite set called alphabet
- its elements are called characters or letters
- $|\Sigma|$  is the size of the alphabet (number of different characters)
- a string over  $\Sigma$  is a finite sequence of characters from  $\Sigma$

- Σ a finite set called alphabet
- its elements are called characters or letters
- $|\Sigma|$  is the size of the alphabet (number of different characters)
- a string over  $\Sigma$  is a finite sequence of characters from  $\Sigma$
- we write strings as  $s = s_1 s_2 \dots s_n$  i.e.  $s_i$  is the *i*'th character of s

- Σ a finite set called alphabet
- its elements are called characters or letters
- $|\Sigma|$  is the size of the alphabet (number of different characters)
- a string over  $\Sigma$  is a finite sequence of characters from  $\Sigma$
- we write strings as  $s = s_1 s_2 \dots s_n$  i.e.  $s_i$  is the i'th character of s

**N.B.**: We number strings from 1, not from 0

• |s| is the length of string s

- |s| is the length of string s
- $\epsilon$  is the empty string, the (unique) string of length 0

- |s| is the length of string s
- $\epsilon$  is the empty string, the (unique) string of length 0
- $\Sigma^n$  is the set of strings of length n

- |s| is the length of string s
- $\epsilon$  is the empty string, the (unique) string of length 0
- $\Sigma^n$  is the set of strings of length n
- $\Sigma^* = \bigcup_{n=0}^{\infty} \Sigma^n$

- |s| is the length of string s
- $\epsilon$  is the empty string, the (unique) string of length 0
- $\Sigma^n$  is the set of strings of length n
- $\Sigma^* = \bigcup_{n=0}^{\infty} \Sigma^n = \Sigma^0 \cup \Sigma^1 \cup \Sigma^2 \cup \dots$  is the set of all strings over  $\Sigma$

#### **Examples**

• DNA:  $\Sigma = \{A,C,G,T\}$ , alphabet size  $|\Sigma| = 4$ , s = ACCTG is a string of length 5 of  $\Sigma$ , with  $s_1 = A, s_2 = s_3 = C, s_4 = T, s_5 = G.$ 

#### **Examples**

- DNA:  $\Sigma = \{A,C,G,T\}$ , alphabet size  $|\Sigma| = 4$ , s = ACCTG is a string of length 5 of  $\Sigma$ , with  $s_1 = A, s_2 = s_3 = C, s_4 = T, s_5 = G$ .
- RNA:  $\Sigma = \{A,C,G,U\}$ , again alphabet size is 4

#### **Examples**

- DNA:  $\Sigma = \{A,C,G,T\}$ , alphabet size  $|\Sigma| = 4$ , s = ACCTG is a string of length 5 of  $\Sigma$ , with  $s_1 = A$ ,  $s_2 = s_3 = C$ ,  $s_4 = T$ ,  $s_5 = G$ .
- RNA: Σ = {A,C,G,U}, again alphabet size is 4
- protein:  $\Sigma = \{A,C,D,E,F,...,W,Y\}$ , alphabet size is 20, ANRFYWNL is a string over  $\Sigma$  of length 8

#### **Examples**

- DNA:  $\Sigma = \{A,C,G,T\}$ , alphabet size  $|\Sigma| = 4$ , s = ACCTG is a string of length 5 of  $\Sigma$ , with  $s_1 = A$ ,  $s_2 = s_3 = C$ ,  $s_4 = T$ ,  $s_5 = G$ .
- RNA: Σ = {A,C,G,U}, again alphabet size is 4
- protein: Σ = {A,C,D,E,F,...,W,Y}, alphabet size is 20,
  ANRFYWNL is a string over Σ of length 8
- English alphabet:  $\Sigma = \{a,b,c,\ldots,x,y,z\}$  of size 26, alphabet is a string over  $\Sigma$  of length 8

Let 
$$s = s_1 \dots s_n$$
 be a string over  $\Sigma$ .

Let 
$$s = s_1 \dots s_n$$
 be a string over  $\Sigma$ .

ex. 
$$s = ACCTG$$

• t is a substring of s if  $t = \epsilon$  or  $t = s_i \dots s_j$  for some  $1 \le i \le j \le n$  (i.e., a "contiguous piece" of s)

Let 
$$s = s_1 \dots s_n$$
 be a string over  $\Sigma$ .

ex. 
$$s = ACCTG$$

• t is a substring of s if  $t = \epsilon$  or  $t = s_i \dots s_j$  for some  $1 \le i \le j \le n$  (i.e., a "contiguous piece" of s)

Let  $s = s_1 \dots s_n$  be a string over  $\Sigma$ .

- t is a substring of s if  $t = \epsilon$  or  $t = s_i \dots s_j$  for some  $1 \le i \le j \le n$  (i.e., a "contiguous piece" of s)
- t is a prefix of s if  $t = \epsilon$  or  $t = s_1 \dots s_j$  for some  $1 \le j \le n$  (i.e., a "beginning" of s)

Let  $s = s_1 \dots s_n$  be a string over  $\Sigma$ .

```
ex. s = ACCTG
```

- t is a substring of s if  $t = \epsilon$  or  $t = s_i \dots s_j$  for some  $1 \le i \le j \le n$  (i.e., a "contiguous piece" of s) CCT, AC, ...
- t is a prefix of s if  $t = \epsilon$  or  $t = s_1 \dots s_j$  for some  $1 \le j \le n$  (i.e., a "beginning" of s) AC, ACCTG, ...

Let  $s = s_1 \dots s_n$  be a string over  $\Sigma$ .

- t is a substring of s if  $t = \epsilon$  or  $t = s_i \dots s_j$  for some  $1 \le i \le j \le n$  (i.e., a "contiguous piece" of s)
- t is a prefix of s if  $t=\epsilon$  or  $t=s_1\dots s_j$  for some  $1\leq j\leq n$  (i.e., a "beginning" of s) AC, ACCTG,...
- t is a suffix of s if  $t = \epsilon$  or  $t = s_i \dots s_n$  for some  $1 \le i \le n$  (i.e., an "end" of s)

Let  $s = s_1 \dots s_n$  be a string over  $\Sigma$ .

- t is a substring of s if  $t = \epsilon$  or  $t = s_i \dots s_j$  for some  $1 \le i \le j \le n$  (i.e., a "contiguous piece" of s) CCT, AC, ...
- t is a prefix of s if  $t=\epsilon$  or  $t=s_1\dots s_j$  for some  $1\leq j\leq n$  (i.e., a "beginning" of s) AC, ACCTG,...
- t is a suffix of s if  $t = \epsilon$  or  $t = s_i \dots s_n$  for some  $1 \le i \le n$  (i.e., an "end" of s) CCTG, G, ...

Let  $s = s_1 \dots s_n$  be a string over  $\Sigma$ .

- t is a substring of s if  $t = \epsilon$  or  $t = s_i \dots s_j$  for some  $1 \le i \le j \le n$  (i.e., a "contiguous piece" of s) CCT, AC, ...
- t is a prefix of s if  $t=\epsilon$  or  $t=s_1\dots s_j$  for some  $1\leq j\leq n$  (i.e., a "beginning" of s) AC, ACCTG,...
- t is a suffix of s if  $t = \epsilon$  or  $t = s_i \dots s_n$  for some  $1 \le i \le n$  (i.e., an "end" of s)
- t is a subsequence of s if t can be obtained from s by deleting some (possibly 0, possibly all) characters from s

Let  $s = s_1 \dots s_n$  be a string over  $\Sigma$ .

- t is a substring of s if  $t = \epsilon$  or  $t = s_i \dots s_j$  for some  $1 \le i \le j \le n$  (i.e., a "contiguous piece" of s)
- t is a prefix of s if  $t=\epsilon$  or  $t=s_1\dots s_j$  for some  $1\leq j\leq n$  (i.e., a "beginning" of s) AC, ACCTG,...
- t is a suffix of s if  $t = \epsilon$  or  $t = s_i \dots s_n$  for some  $1 \le i \le n$  (i.e., an "end" of s) CCTG, G, ...
- t is a subsequence of s if t can be obtained from s by deleting some
   (possibly 0, possibly all) characters from s
   AT, CCT,...

Let  $s = s_1 \dots s_n$  be a string over  $\Sigma$ .

**ex.** s = ACCTG

- t is a substring of s if  $t = \epsilon$  or  $t = s_i \dots s_j$  for some  $1 \le i \le j \le n$  (i.e., a "contiguous piece" of s)
- t is a prefix of s if  $t=\epsilon$  or  $t=s_1\dots s_j$  for some  $1\leq j\leq n$  (i.e., a "beginning" of s) AC, ACCTG, ...
- t is a suffix of s if  $t = \epsilon$  or  $t = s_i \dots s_n$  for some  $1 \le i \le n$  (i.e., an "end" of s)

#### N.B.

string = sequence, but substring  $\neq$  subsequence!

#### N.B.

1. Every substring is a subsequence, but not every subsequence is a substring!

#### N.B.

1. Every substring is a subsequence, but not every subsequence is a substring!

**Ex.:** Let s = ACCTG, then ACT is a subsequence but not a substring.

#### N.B.

1. Every substring is a subsequence, but not every subsequence is a substring!

**Ex.:** Let s = ACCTG, then ACT is a subsequence but not a substring.

2. Every prefix and every suffix is a substring.

#### N.B.

- 1. Every substring is a subsequence, but not every subsequence is a substring!
  - **Ex.:** Let s = ACCTG, then ACT is a subsequence but not a substring.
- 2. Every prefix and every suffix is a substring.
- 3. t is substring of  $s \Leftrightarrow t$  is prefix of a suffix of  $s \Leftrightarrow t$  is suffix of a prefix of s

# Counting substrings, subsequences etc.

#### Question

Given  $s = s_1 \dots s_n$ . How many

- prefixes,
- suffixes,
- substrings,
- subsequences

does s have (exactly, or at most, or at least)?