# In The Name of God



# Simple and Efficient Pattern Matching Algorithms for Biological Sequences

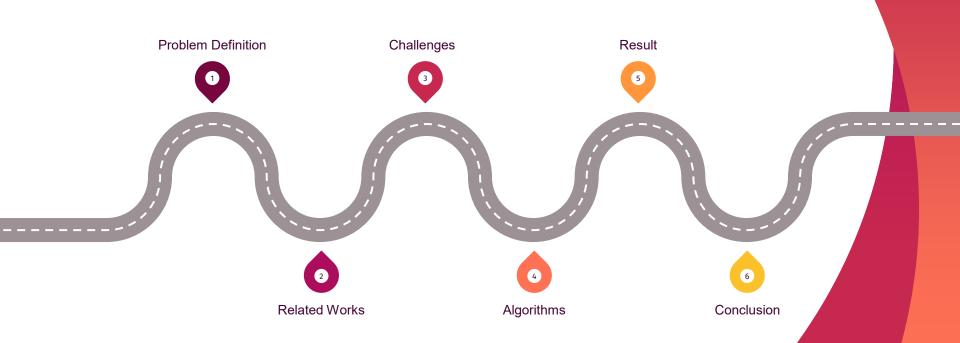
#### **Presenters:**

Nazanin Fereydoonizade – Sarvin Saravi

### **Suppervisor:**

Dr. Shakibian

# **Road Map**



# What is Pattern Matching?

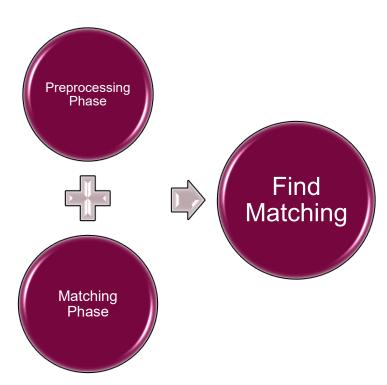
Scan a sequence or a database to detect the locations of a pattern

## **Applications**

- 1. Image & Signal Processing
- 2. Information Retrieval
- 3. Search Engine
- Text Processing
- Parsers
- 6. NLP



# **Pattern Matching Phases**



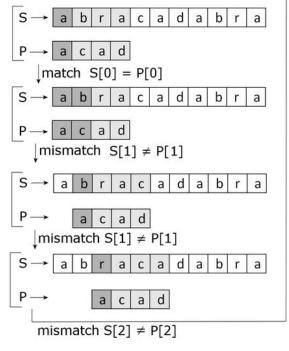
## **Related Works**

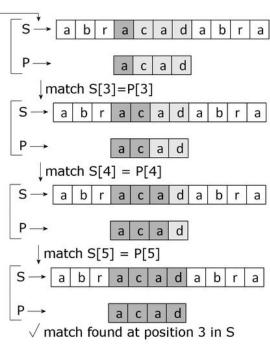
- 1. Brute Force Time problem, O(mn)
- 2. **DFA-bases** Not Scalable, O(n)
- 3. KMP Not Suitable for Small Strings, O(n)
- 4. Boyer Moore Depend on String size, O(m+n) /
- **5. DCPM** O(mn)

Related

Works

## 1. Brute Force





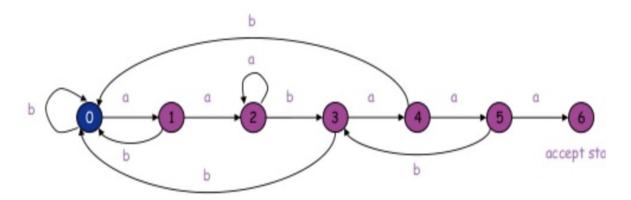
## 2. DFA-based

- Build DFA from Pattern
- Run DFA on Text

### example

Text: aaabaabaaab

Pattern: aabaaa



# Challenges

- 1. Library
- 2. Optimal File Size
- 3. Execution Time
- 4. Performance Analysis Using Different Encoding Techniques

# Algorithm.1: First-Last Pattern Matching (Preprocessing Phase)

### **Algorithm 1** Preprocessing Phase of FLPM Algorithm

**Input:** Text t and pattern p stored in the arrays of t[0..n-1] and p[0..m-1], respectively, over finite alphabet  $\sum$ .

**Output:** The number of windows identified in this phase and their start indexes.

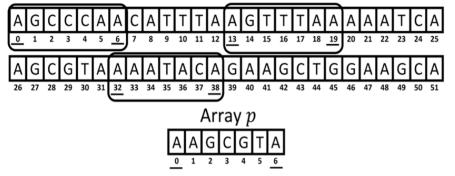
- 1.  $count \leftarrow 0$
- 2.  $num\_window \leftarrow 0$
- 3. WHILE  $count \le n m$  DO
- 4. **IF** t[count] = p[0] **THEN**
- 5. **IF** t[count + m 1] = p[m 1] **THEN**
- 6.  $window\_index[num\_window] \leftarrow count$
- 7.  $num\_window \leftarrow num\_window + 1$
- 8. **END-IF**
- 9. END-IF
- 10.  $count \leftarrow count + 1$
- 11. END-WHILE

# Algorithm.1: First-Last Pattern Matching (Matching Phase)

```
Algorithm 2 Matching Phase of FLPM Algorithm
  Input: The number of windows identified in the
        preprocessing phase and their start indexes.
  Output: The start index for all occurrences of pattern p in
        text t.
  1. count \leftarrow 0
  2. num_match \leftarrow
  3. WHILE count < num_window DO
           s \leftarrow window\_index[count]
           c \leftarrow 1
  6.
           WHILE c < m - 2 DO
                IF p[c] \neq t[s+c] THEN
  8.
                      BREAK /*Exit the current loop*/
                END-IF
 10.
                c \leftarrow c + 1
 11.
          END-WHILE
 12.
          IF c = m - 1 THEN
 13.
                match\_index[num\_match] \leftarrow s
 14.
                num\ match \leftarrow num\ match + 1
 15.
          END-IF
 16.
          count \leftarrow count + 1
 17. END-WHILE
```

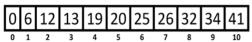
# Algorithm.1: First-Last Pattern Matching (Example)

### Array t



Array window\_index

Array *match\_index* 





# Algorithm.2: Processor-Aware Pattern Matching (Preprocessing Phase)

#### **Algorithm 3** PAPM Algorithm

```
Input: Text t and pattern p stored in the arrays of t[0..n-1] and p[0..m-1], respectively, over finite alphabet \sum.
```

**Output:** The start index for all occurrences of pattern p in text t.

#### /\* PREPROCESSING PHASE \*/

- 1.  $count \leftarrow 0$
- 2.  $num\_window \leftarrow 0$
- 3. WHILE  $count \le n m$  DO
- 4. **IF** t [count..count + word\_len 1]
- 5.  $= p[0..word\_len 1]$  **THEN**
- 6.  $window_index[num\_window] \leftarrow count$
- 7.  $num\_window \leftarrow num\_window + 1$
- 8. **END-IF**
- 9.  $count \leftarrow count + 1$
- 10. END-WHILE

# Algorithm.2: Processor-Aware Pattern Matching (Matching Phase)

```
11. k \leftarrow m \mod word\_len
12. IF k = 0 THEN
13.
           start\_index \leftarrow word\_len
14. ELSE
15.
           start\_index \leftarrow k
16. END-IF
17. count \leftarrow 0
18. num\_match \leftarrow 0
19. WHILE count < num_window
                                         DO
20.
          s \leftarrow window\_index[count]
21.
          c \leftarrow start\_index
22.
          WHILE c < m - word len
23.
              IF p[c..c + word\_len - 1]
24.
                \neq t[s+c..s+c+word\_len-1] THEN
25.
                         BREAK
26.
              END-IF
27.
              c \leftarrow c + word \ len
28.
           END-WHILE
29.
           IF c = m THEN
30.
               match\_index[num\_match] \leftarrow s
31.
               num\_match \leftarrow num\_match + 1
32.
            END-IF
33.
            count \leftarrow count + 1
34. END-WHILE
```

# Algorithm.2: Processor-Aware Pattern Matching

(Example)

Array t

Array p A A G C G T A

Array window\_index Array match\_index





# Algorithm.3: Least frequency pattern matching algorithm

- LFPM
- an enhancement of PAPM
- specialized for DNA
- many patterns vs. few patterns

# Least frequency pattern matching algorithm

The first step before the preprocessing phase

- $\Sigma = \{ A, C, G, T \} \longrightarrow HRG$
- word\_len = b/8 for b-bit computer
- all possible words = 4<sup>word\_len</sup>
- Create freq\_table

# Least frequency pattern matching algorithm

The first step before the preprocessing phase

**TABLE 1. THE freq\_table.** 

	0	1	2	3	4	5		$word\_len-2$	$word\_len-1$	word_len
0	Α	Α	Α	Α	Α	Α		А	Α	$n_0$
1	Α	Α	Α	Α	Α	Α		А	С	$n_1$
2	Α	Α	Α	Α	Α	Α		А	G	$n_2$
3	Α	Α	Α	Α	Α	Α		А	Т	$n_3$
$4^{word\_len} - 2$	Т	Т	Т	Т	Т	Т		Т	G	$n_{4^{word\_len}-2}$
$4^{word\_len} - 1$	Т	Т	Т	Т	Т	Т		Т	Т	$n_{4^{word\_len}-1}$

#### Algorithm 4 Filling the Table of Frequency

**Input:** The array of reference and the length of this array. **Output:** The filling table of word frequency.

- 1.  $freq\_table[0..4^{word\_len} 1][0..word\_len 1] \leftarrow$
- 2. All possible words with word\_len length over  $\Sigma = \{ACGT\}$
- 3.  $freq\_table[0..4^{word\_len} 1][word\_len] \leftarrow 0$
- 4.  $count \leftarrow 0$
- 5. WHILE  $count \le REF\_len word\_len$  **DO**
- 6.  $row\_count \leftarrow 0, col\_count \leftarrow 0$
- 7.  $w[0..word\_len 1] \leftarrow REF$  $[count..count + word\_len - 1]$
- 8. WHILE col\_count < word\_len DO
- 9. **SWITCH** w [col\_count]
- 10. CASE 'A': nothing
- 11. CASE 'C': Increase  $row\_count$  by  $1 \times 4^{word\_len-col\_count-1}$
- 12. CASE 'G': Increase  $row\_count$  by  $2 \times 4^{word\_len-col\_count-1}$
- 13. **CASE** 'T': Increase  $row\_count$  by  $3 \times 4^{word\_len-col\_count-1}$
- 14. END- SWITCH
- 15.  $col\_count \leftarrow col\_count + 1$
- 16. END-WHILE
- Increase freq\_table[row\_count][word\_len] by one
- 18.  $count \leftarrow count + 1$
- 19. END-WHILE

# Least frequency pattern matching algorithm

The preprocessing & Matching phase

words in the pattern → least frequent word → start index

similar to PAPM

# Least frequency pattern matching algorithm

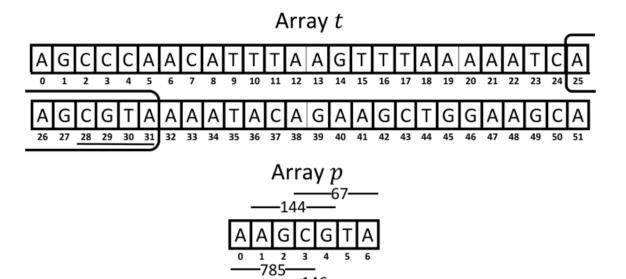


FIGURE 3. An example for the operation of LFPM algorithm.

#### Algorithm 5 LFPM Algorithm

**Input:** The filled  $freq\_table$  on reference data. Text t and pattern p stored in the arrays of t[0..n-1] and p[0..m-1], respectively, over finite alphabet  $\sum = \{A, C, G, T\}$ . **Output:** The first indexes for all occurrences of pattern p in text t.

#### /\*PREPROCESSING PHASE\*/

## /\*Step 1: finding the least frequent word in the pattern\*/

- 1.  $count \leftarrow 0$
- 2.  $min_value \leftarrow \infty$
- 3.  $min\_index \leftarrow -1$
- 4. WHILE  $count \le m word\_len$  **DO**
- 5.  $row\_count \leftarrow 0, col\_count \leftarrow 0$
- 6.  $w[0..word\_len 1] \leftarrow p[count..count + word\_len 1]$
- WHILE col\_count < word\_len DO</li>
- 8. **SWITCH** w [col\_count]
- CASE'A': nothing
- CASE'C': Increase row\_count by 1 × 4<sup>word\_len-col\_count-1</sup>
- 11. CASE'G': Increase row\_count by 2 × 4word\_len-col\_count-1
- 12. **CASE**'T': Increase row\_count by  $3 \times 4^{word\_len-col\_count-1}$

```
13. END- SWITCH
```

- 14.  $col\_count \leftarrow col\_count + 1$
- 15. END-WHILE
- 16. **IF** freq\_table[row\_count][word\_len] < min\_value **THEN**
- 17.  $min\_value=freq\_table[row\_count][word\_len]$
- 18.  $min\_index \leftarrow count$
- 19. **END-IF**
- 20.  $count \leftarrow count + 1$
- 21. END-WHILE

Time complexity:  $\theta(\text{word\_len} \times (\text{m - word\_len})) \approx \theta(\text{m})$ 

### /\*preprocessing phase\*/

### /\*Step 2: finding the windows\*/

- 22.  $count \leftarrow min\_index$
- 23.  $num\_window \leftarrow 0$
- 24. WHILE count  $\leq n (m min\_index)$  **DO**
- 25. **IF** t [count..count + word\_len 1]
- 26. =  $p[min\_index..min\_index + word\_len 1]$ **THEN**
- 27. window\_index[num\_window] ← count min\_index
- 28.  $num\_window \leftarrow num\_window + 1$
- END-IF
- 30.  $count \leftarrow count + 1$
- 31. END-WHILE

#### /\*MATCHING PHASE\*/

```
32. count \leftarrow 0
33. num\_match \leftarrow 0
34. k \leftarrow mmodword\_len
35. WHILE count < num_window DO
36.
            s \leftarrow window\_index[count]
37.
            c \leftarrow 0
38.
            w \leftarrow word\_len
            WHILE c \le m - 1 DO
39.
40.
                IF c > m - word\_len THEN
41.
                       w \leftarrow k
42.
                END-IF
43.
                IF p[c..c + w - 1]
44.
                  \neq t[s + c...s + c + w - 1] THEN
45.
                      BREAK
                END-IF
46.
47.
               c \leftarrow c + w
            END-WHILE
48.
            IF c = m THEN
49.
50.
                match\_index[num\_match] \leftarrow s
51.
                num_match \leftarrow num_match + 1
            END-IF
52.
53.
            count \leftarrow count + 1
54. END-WHILE
```

- ☐ Brute Force (BF)
- Boyer-Moore (BM)
- Divide and Conquer Pattern Matching (DCPM)

FLPM

PAPM

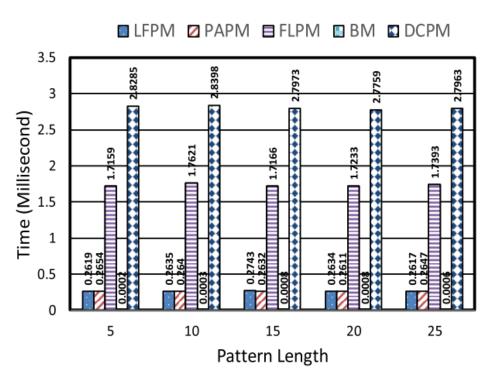
■ LFPM

The specifications of the computing environment

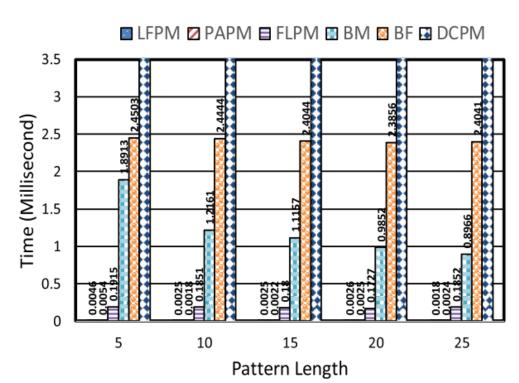
Intell®core<sup>TM</sup>2 Duo CPU T6600 (a 2.2 GHz clock) A 2GB Memory Acer (Aspire 5738)

Windows 7 ultimate 32 bit

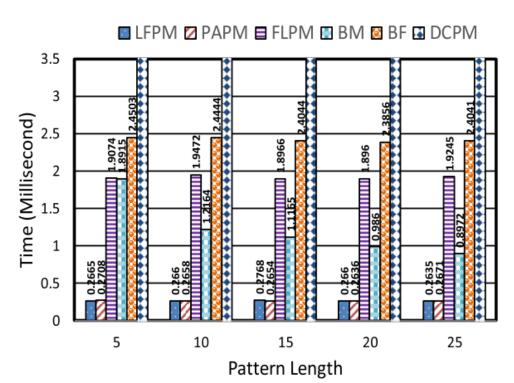
### A. THE TIME OF PREPROCESSING PHASE



### B. THE TIME OF MATCHING PHASE



### C. TOTAL TIME OF PATTERNS

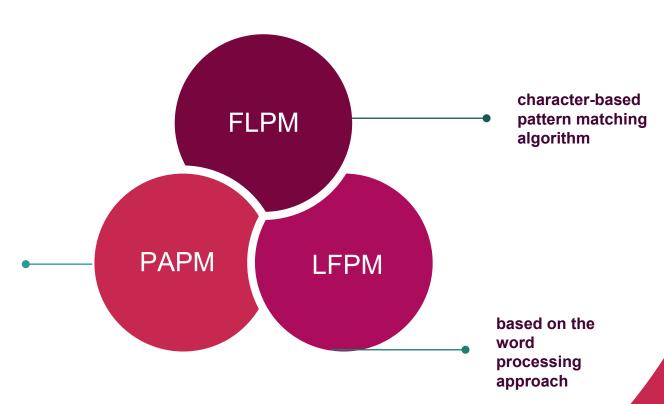


based on the

processing approach

word

## **Conclusion**



### **Future Research**

- A parallel version of the presented algorithms
- The algorithms supporting approximate matching

### Reference

- Neamatollahi, Peyman, Montassir Hadi, and Mahmoud Naghibzadeh. "Simple and efficient pattern matching algorithms for biological sequences." IEEE Access 8 (2020): 23838-23846.
- > SAQIB HAKAK1, Amirrudin Kamsin1, PALAIAHNAKOTE SHIVAKUMARA1, GULSHIN "Exact String Matching Algorithms: Survey, Issues, and Future Research Directions"
- https://en.wikipedia.org/wiki/Boyer-Moore\_stringsearch\_algorithm
- https://en.wikipedia.org/wiki/Knuth-Morris-Pratt\_algorithm
- https://en.wikipedia.org/wiki/Brute-force\_attack

