### CMSC423: Bioinformatic Algorithms, Databases and Tools

Exact string matching: introduction

### Sequence alignment: exact matching

```
Text
CAGGTACAGTTCCCTCGACACCTACTACCTAAG
                                                      Pattern
 СТАСТ
 ССТАСТ
  CCTACT
      for i = 0 .. len(Text) {
       for j = 0 .. len(Pattern) {
         if (Pattern[i] != Text[i]) go to next i
       if we got there pattern matches at i in Text
```

```
Running time = O(len(Text) * len(Pattern)) = O(mn)
```

What string achieves worst case?

#### Worst case?

(m - n + 1) \* n comparisons

#### Can we do better?

the Z algorithm (Gusfield)

For a string T, Z[i] is the length of the longest prefix of T[i..m] that matches a prefix of T. Z[i] = 0 if the prefixes don't match.

```
T[0 ... Z[i]] = T[i ... i+Z[i] -1]
```

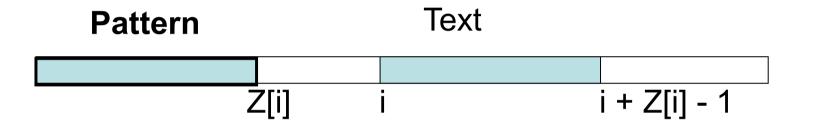


#### Example Z values

#### ACAGGTACAGTTCCCTCGACACCTACTACCTAG 00100040100000000302000200200110

## Can the Z values help in matching?

Create string Pattern\$Text where \$ is not in the alphabet



If there exists i, s.t. Z[i] = length(Pattern) Pattern occurs in the Text starting at i

#### example matching

CCTACT\$ACAGGTACAGTTCCCTCGACACCTACTACCTAG 0100100010000100002310100106100100410000

• What is the largest Z value possible?

# Can Z values be computed in linear time?

Z[1]? compare T[1] with T[0], T[2] with T[1], etc. until mismatch Z[1] = 2

This simple process is still expensive:

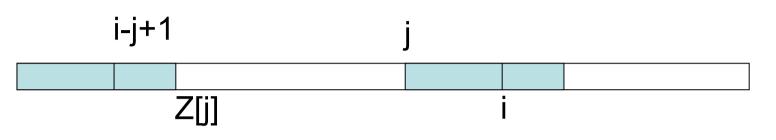
T[2] is compared when computing both Z[1] and Z[2].

Trick to computing Z values in linear time: each comparison must involve a character that was not compared before

Since there are only m characters in the string, the overall # of comparisons will be O(m).

# Basic idea: 1-D dynamic programming

Can Z[i] be computed with the help of Z[j] for j < i?

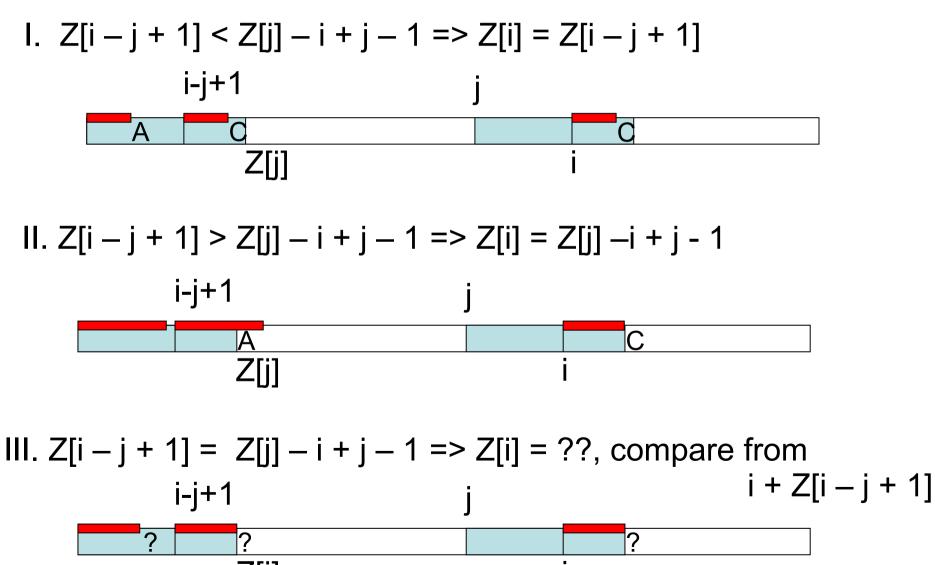


Assume there exists j < i, s.t. j + Z[j] - 1 > ithen Z[i - j + 1] provides information about Z[i]

If there is no such j, simply compare characters T[i..] to T[0..] since they have not been seen before.

#### Three cases

Let j < i be the coordinate that maximizes j + Z[j] - 1 (intuitively, the Z[j] that extends the furthest)



# Time complexity analysis

- Why do these tricks save us time?
- 1. Cases I and II take constant time per Z-value computed total time spent in these cases is O(n)
- 2. Case III might involve 1 or more comparisons per Z-value however:

- every successful comparison (match) shifts the rightmost character that has been visited

- every unsuccessful comparison terminates the "round" and algorithm moves on to the next Z-value

total time spent in III cannot be more than # of characters in the text

Overall running time is O(n) CMSC423 Fall 2009

# Space complexity?

 If using Z algorithm for matching, how many Z values do we need to store?

#### 

 Only need to remember Z-values for pattern and the "farthest reaching Z-value" (Z[j] in what we discussed before)

## Some questions

• What are the Z-values for the following string:

TTAGGATAGCCATTAGCCTCATTAGGGATTAGGAT

- In the string above, what is the longest prefix that is repeated somewhere else in the string?
- Trace through the execution of the linear-time algorithm for computing the Z values for the string listed above. How many times do rules I, II, and III apply?

# Z algorithm, not just for matching

• Lempel-Ziv compression (e.g. gzip)

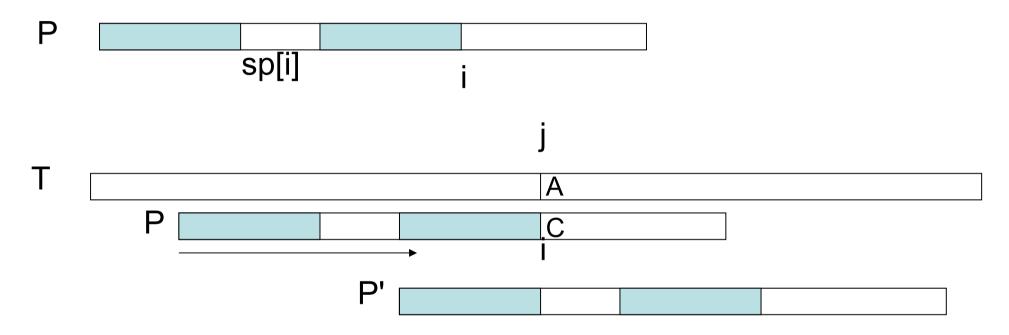


if Z[i] = 0, just send/store the character T[i], otherwise, instead of sending T[i..i+Z[i] - 1] (Z[i] - 1 characters/bytes) simply send Z[i] (one number)

 Note: other exact matching algorithms used for data compression (e.g. Burrows-Wheeler transform relates to suffix arrays)

# Knuth-Morris-Pratt algorithm

Given a Pattern and a Text, preprocess the Pattern to compute sp[i] = length of longest prefix of P that matches a suffix of P[0..i]

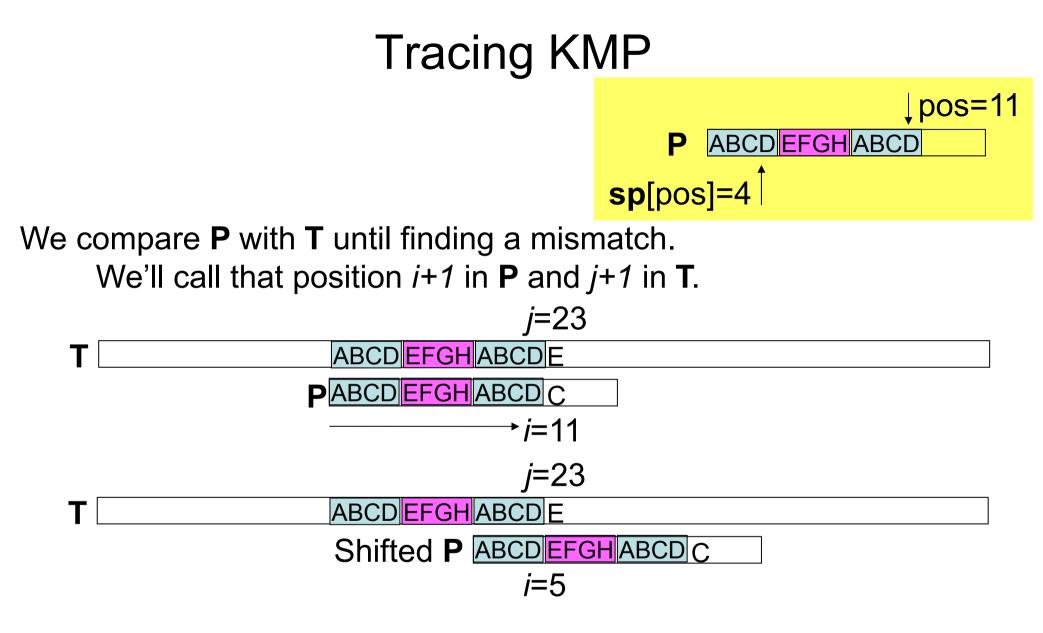


- Compare P with T until finding a mis-match (at coordinate i + 1 in P and j + 1 in T).
- Shift P such that first sp[i] characters match T[j sp[i] + 1 .. j].
- Continue matching from T[i+1], P[sp[i]+1]

# Knuth-Morris-Pratt (KMP) Algorithm

Given a pattern (**P**) and a text block (**T**) you preprocess **P** to compute a zero-indexed array **sp**[] where **sp**[pos] contains the length of longest **prefix** of **P** that matches a **suffix** of **P**[0..pos]

Next 4 slides from Evan Golub



We then logically shift **P** using the **sp** value.

This allows the first **sp**[*i*] characters to match **T**[(*j*-sp[*i*]+1) .. *j*]. We then continue comparing from **P**[sp[*i*]+1] and **T**[*j*+1] CMSC423 Fall 2009

index: 0123456

- pattern: AAAAAAA
- sp: 0123456

index: 0123456

- pattern: AAAAAB
- sp: 0123450

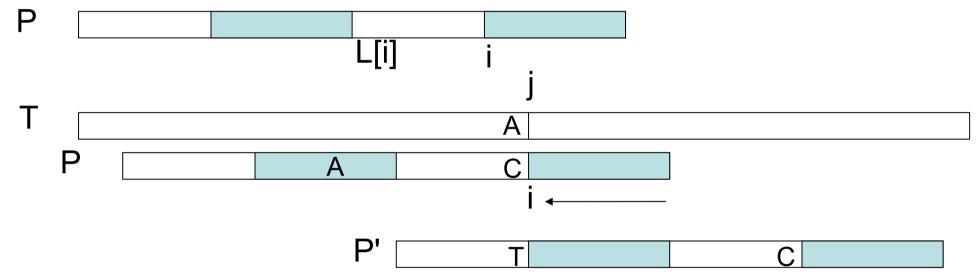
#### AAAABAAAAABAAAAAAA

- index: 0123456
- pattern: ABACABC
- sp: 0010120

#### ABABBABAABABACABC

# **Boyer-Moore algorithm**

Preprocess the pattern, computing, for every i, L[i] = largest coordinate < n, s.t. P[i..n] matches a suffix of P[1..L[i]] (inverted Z function)

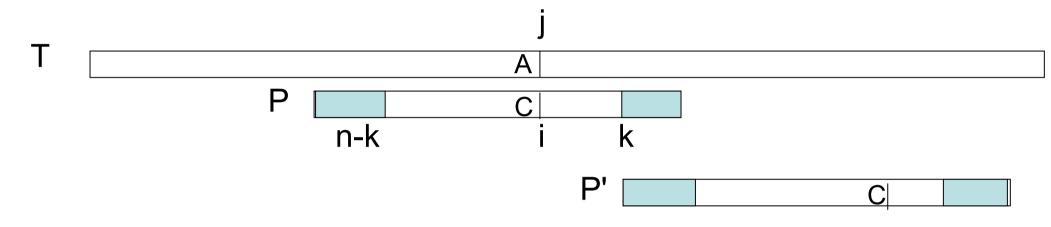


Match the pattern backwards (starting at the right) until mismatch. Shift the pattern such that P[L[i] - n + i + 1] matches at T[j]Repeat.

Bad character rule: find character T[j - 1] in P and shift until it matches. Choose the longest shift (btwn. suffix & char. rules)

# Boyer-Moore ... cont

- What if P[i..n] does not occur elsewhere prior to i?
- Find k > i s.t. P[1..(n-k)] matches P[k..n]



- Also bad character rule:
  - if P[i] mismatches with T[j] can shift P until we find a character equal to T[j] in P (above, shift until an A in P lines up to the A in T)
- Putting it all together: compute the shift according to the suffix rules and the bad character rule and pick the largest

# Questions

- Can you use the Z-values to efficiently compute the sp() values used in the KMP algorithm?
- How about the values used by the Boyer-Moore algorithm?