

# String Algorithms and Data Structures

## Z-values and the Z-algorithm

CS 199-225

February 6, 2023

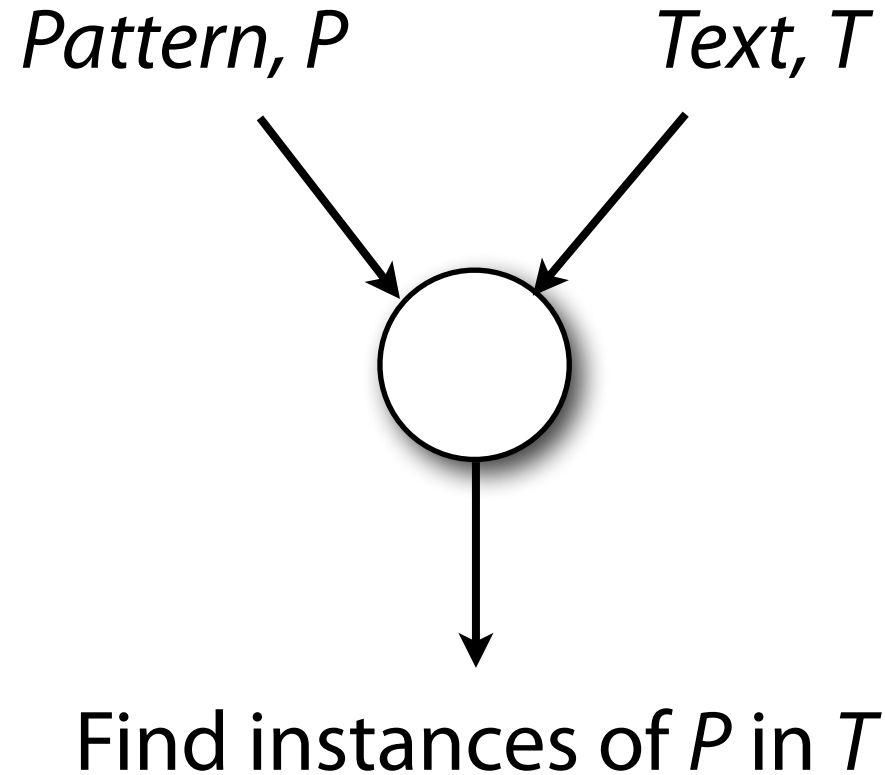
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# Exact Pattern Matching



'instances': An exact, full length copy

# Exact Pattern Matching

What's a simple algorithm for exact matching?

P: word

T: There would have been a time for such a word

word word word word word word word word word **word**

word word word word word word word word

word word word word word word word word

word word word word word word word word

word word word word word word word word

← One occurrence

Try all possible alignments. For each, check if it matches. This is the *naïve algorithm*.

# Exact Pattern Matching

What is good about the naive solution?

What is bad?

# Exact Pattern Matching

What is our time complexity?

$$(n = |P|, \quad m = |T|)$$

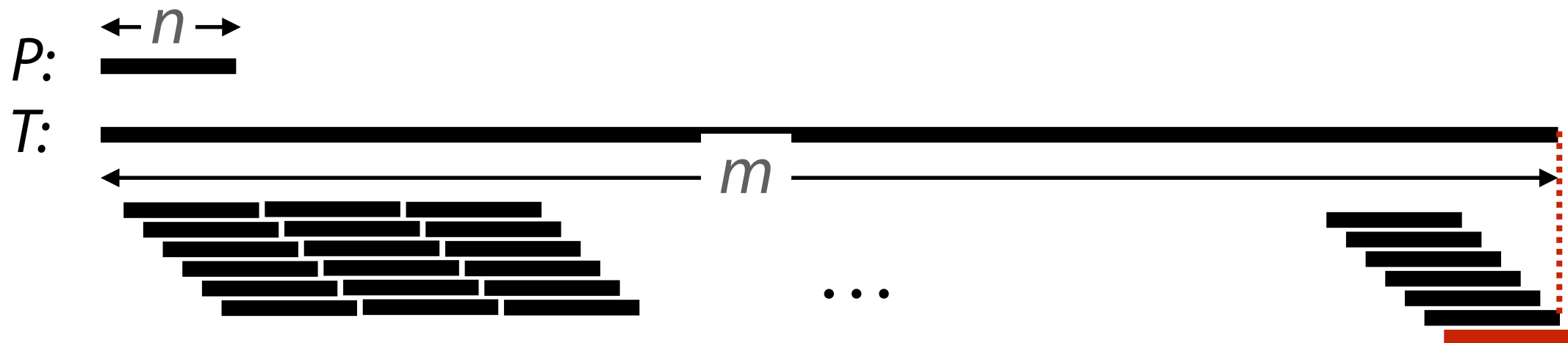
(# of alignments) x (cost of an alignment)

# Exact Pattern Matching

What is our time complexity?

$$(n = |P|, \quad m = |T|)$$

(# of alignments) x (cost of an alignment)



P can fit at each 'position' along T except the **edge**

# Exact Pattern Matching

What is our time complexity?

$$(n = |P|, \quad m = |T|)$$

( \_\_\_\_\_ ) x (cost of an alignment)

*P*: aaaa

*T*: aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa **aaaa**

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa **aaaa**

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa **aaaa**

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa

There are \_\_\_\_\_ positions which extend past the edge of T

# Exact Pattern Matching

What is our time complexity?

$$(n = |P|, \quad m = |T|)$$

$$(m-n+1) \times (\text{cost of an alignment})$$

*P*: aaaa

*T*: aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa

aaaa aaaa aaaa aaaa aaaa aaaa aaaa aaaa

Each alignment compares \_\_\_\_\_ characters.



# Exact Pattern Matching

What is our time complexity?

$(n = |P|, \quad m = |T|)$

$$\theta((m - n + 1) \times n)$$

# String Algorithms in Genomics

P: Read (  $n = \sim 50-150$  )

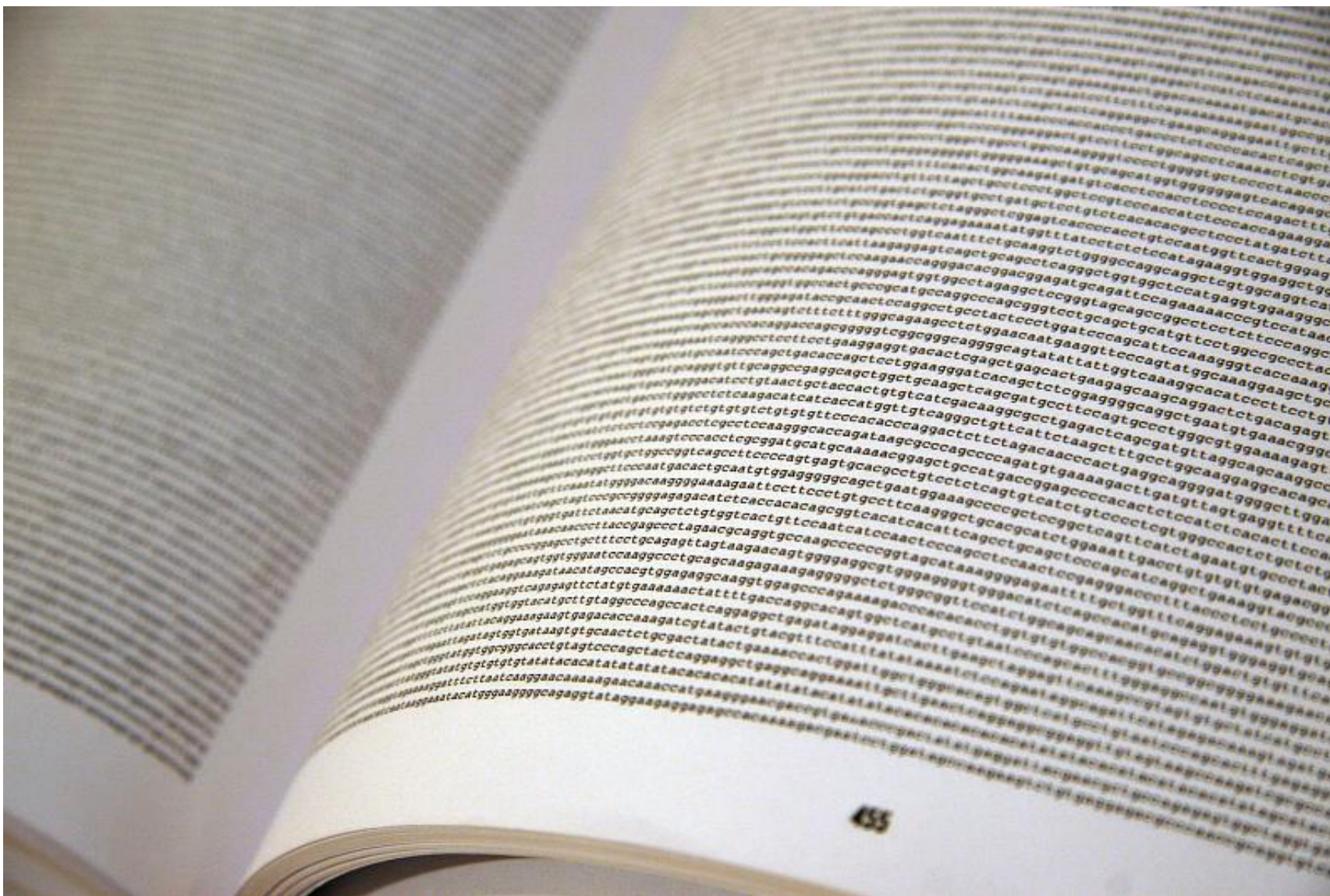
CTCAAACCTCTGACCTTTGGTGATCCACCCGCCTAGGCCTTC

T: Reference (  $m = \sim 3$  billion )

GATCACAGGTCTATCACCTATTAACCACTCACGGGAGCTCTCCATGCATTTGGTATTTT  
CGTCTGGGGGATGCACGCGATAGCATTGCGAGACGCTGGAGCCGGAGCACCTATGTC  
GCAGTATCTGTCTTTGATTCTGCCTCATCTATTATTTATCGCACCTACGTTCAATATT  
ACAGGCGAACATACTTACTAAAGTGTGTTAATTAATTAATGCTTGTAGGACATAATAATA  
ACAATTGAATGTCTGCACAGCCACTTTCCACACAGACATCATAACAAAAATTTCCACCA  
AACCCCCCTCCCCGCTTCTGGCCACAGCACTTAAACACATCTCTGCCAAACCCAAAA  
ACAAAGAACCCTAACACCAGCCTAACAGATTTCAAATTTTATCTTTTGGCGGTATGCAC  
TTTTAACAGTCACCCCCCACTAACACATTATTTTCCCCTCCCCTCCACTCCATACTAAT  
CTCATCAATACAACCCCCGCCATCCTACCCAGCACACACACACCCGCTGCTAACCCATA  
CCCCGAACCAACCAACCCCAAGACACCCCCACAGTTTATGTAGCTTACCTCCTCAA  
GCAATACACTGACCCGCTCAAACCTCTGGATTTTGGATCCACCCAGCGCCTTGGCCTAAA  
CTAGCCTTTCTATTAGCTCTTAGTAAGATTACACATGCAAGCATCCCCGTTCCAGTGAGT  
TCACCTCTAAATCACCACGATCAAAGGAACAAGCATCAAGCACGCAATGCAGCTC  
AAAACGCTTAGCCTAGCCACACCCCCACGGGAAACAGCAGTGATTAACCTTAGCTATAA  
ACGAAAGTTAACTAAGCTATACTAACCCAGGGTTGGTCAATTTGGTCCAGCCACCTC  
GGTCACACGATTAACCCAAGTCAATAGAAGCCGGCGTAAAGAGGTTTGTAGATCACCC  
TCCCCAATAAAGCTAAAACCTCACCTGAGTTGTA AAAA ACTCCGTTGACACAAAATAGAC  
TACGAAAGTGGCTTTAACATATCTGAACACACAATAGCTAAGCCCAA ACTGGGATAGA  
TACCCACTATGCTTAGCCCTAAACCTCAACAGTTAAATCAA AAAACTGCTCGCCAGAA  
CACTACGAGCCACAGCTTAAAACCTCAAAGGACCTGGCGGTGCTCATATCCCTCTAGAGG  
AGCCTGTTCTGTAATCGATAAACCCCGATCAACCTCACCACCTCTGCTCAGCCTATAT  
CCGCCATCTTCAGCAAACCCTGATGAAGGCTACAAAGTAAGCGCAAACCTACCACGTA  
ACGTTAGGTCAAGGTGTAGCCCATGAGGTGGCAAGAAATGGGCTACATTTCTACCCCA  
AAAAC TACGATAGCCCTTATGAAACTTAAGGGTGAAGGTGGATTTAGCAGTAAACTAAG  
AGTAGAGTGCTTAGTTGAACAGGGCCCTGAAGCGGTACACACCCGCCGTCACCTCCTC  
AAGTATACTTCAAAGGACATTTAACTAAAACCCCTACGCATTTATATAGAGGAGACAAGT  
CGTAACCTCAAACCTCTGCCTTTGGTGATCCACCCGCCTTGGCCTACCTGCATAATGAAG  
AAGCACCCTAACCTTACACTTAGGAGATTTCAACTTAACTTACCCCTCTGAGCTAAAGCTA

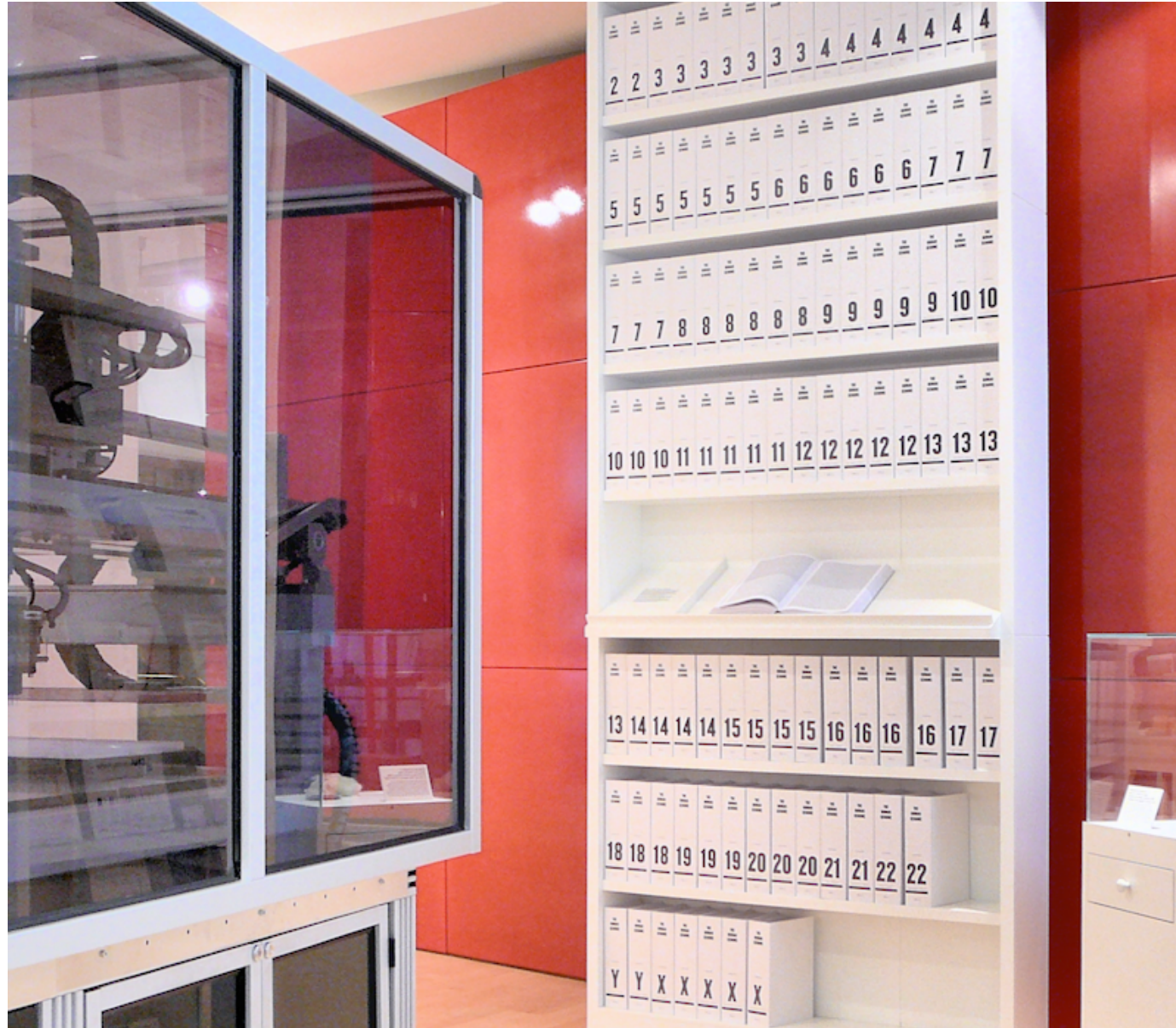


# String Algorithms in Genomics





# String Algorithms in Genomics



# Improving exact pattern matching



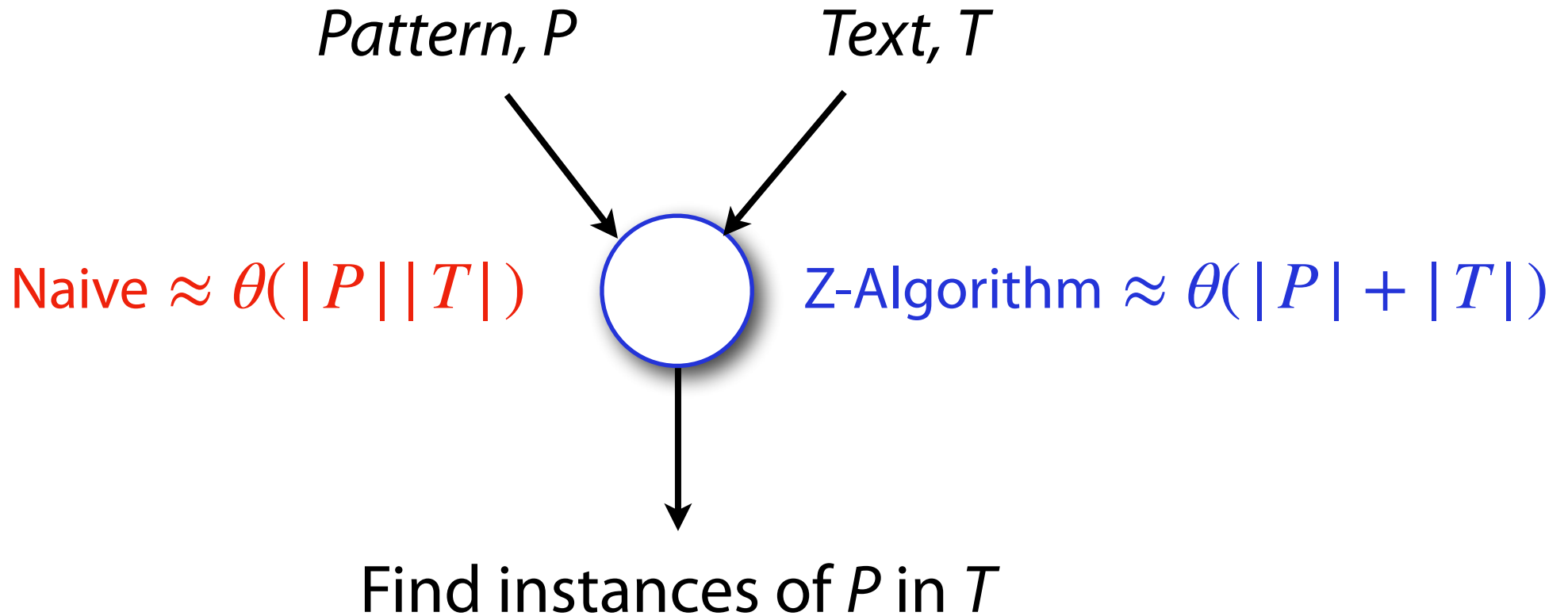
How can we do better than the naïve algorithm?

... If we have infinite space?

... If I tell you the pattern ahead of time?

... If I tell you the text ahead of time?

# Exact Pattern Matching w/ Z-algorithm



'instances': An exact, full length copy

# The Z-value [ $Z_i(S)$ ]

Given a string  $S$ ,  $Z_i(S)$  is the length of the longest substring in  $S$ , starting at position  $i$ , that matches a prefix of  $S$ .

                  0 1 2 3 4 5 6 7 8 9  
S:    **T T C G T T A G C G**

$Z_0(S) =$

$Z_3(S) =$

$Z_1(S) =$

$Z_4(S) =$

$Z_2(S) =$

$Z_5(S) =$

# The Z-value [ $Z_i(S)$ ]

Given a string  $S$ ,  $Z_i(S)$  is the length of the longest substring in  $S$ , starting at position  $i$ , that matches a prefix of  $S$ .

                  0 1 2 3 4 5 6 7 8 9  
S:    **T T C G T T A G C G**

$$Z_0(S) = 10$$

$$Z_1(S) = 1$$

$$Z_2(S) = 0$$

$$Z_3(S) =$$

$$Z_4(S) =$$

$$Z_5(S) =$$



# The Z-value [ $Z_i(S)$ ]

Given a string  $S$ ,  $Z_i(S)$  is the length of the longest substring in  $S$ , starting at position  $i > 0$ , that matches a prefix of  $S$ .

          0 1 2 3 4 5 6 7 8 9  
S:    **T T C G T T A G C G**

$$Z_0(S) = 10$$

$$Z_1(S) = 1$$

$$Z_2(S) = 0$$

$$Z_3(S) = 0$$

$$Z_4(S) = 2$$

$$Z_5(S) = 1$$

# Calculating the Z-values

**Naive:** Compute the Z-values by *explicitly* comparing characters (left-to-right scan):

$$Z_1 =$$

A A A A B A A C A A B A A ...

A A A A B A A C A A B A A ...

$$Z_5 =$$

A A A A B A A C A A B A A ...

A A A A B A A C A A B A A ...

*What is our time complexity?*

# Calculating the Z-values

**Naive:** Compute the Z-values by *explicitly* comparing characters (left-to-right scan):

**S : 1 1 0 1 1 0 0 1**

*What is our time complexity?*

# Calculating the Z-values



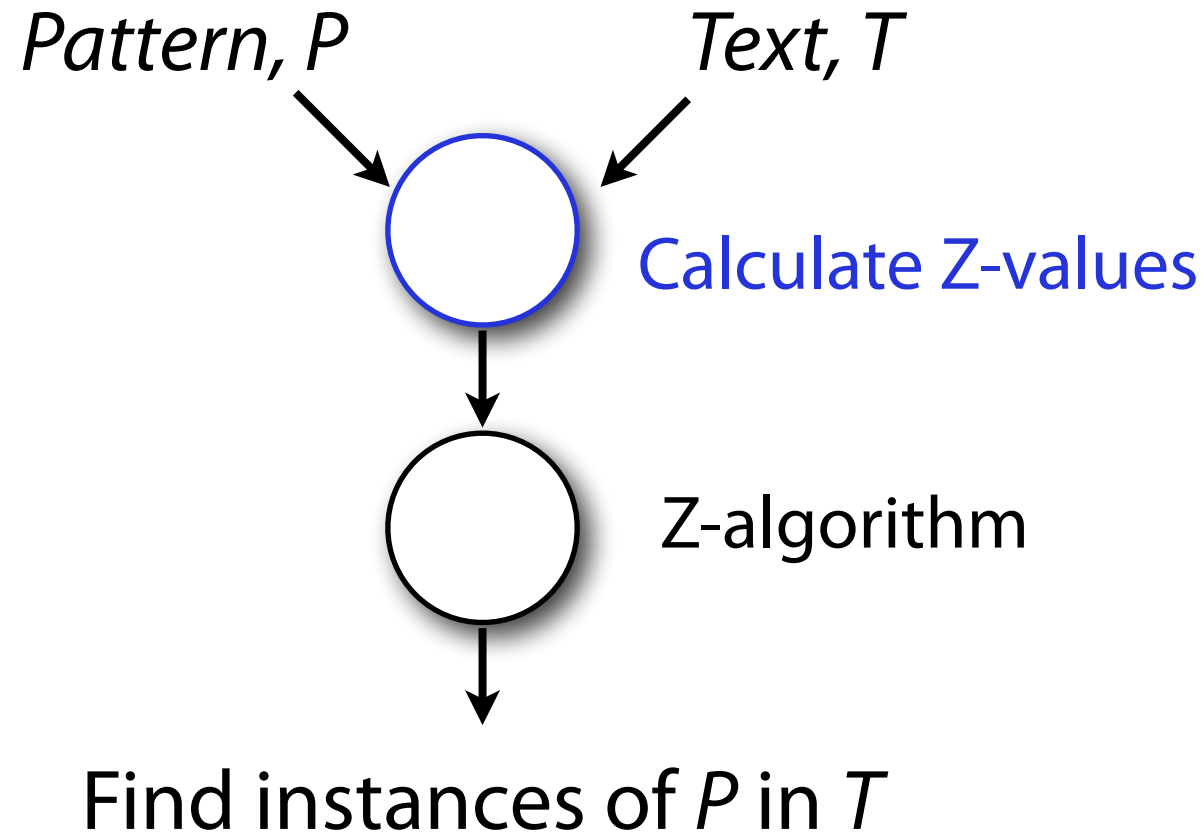
**Naive:** Compute the Z-values by *explicitly* comparing characters (left-to-right scan):

```
S : 1 1 0 1 1 0 0 1
    1 0 1 1 0 0 1
    0 1 1 0 0 1
    1 1 0 0 1
    1 0 0 1
    0 0 1
    0 1
    1
```

*What is our time complexity?*

# Pattern matching with the Z-value

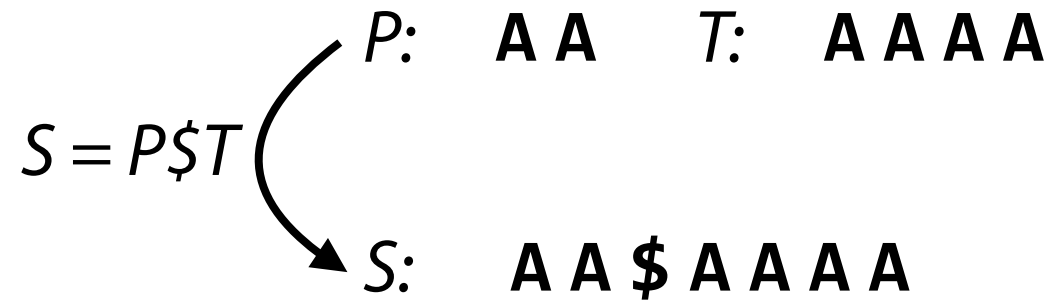
Given a  $Z_i$  value calculator, how do we solve pattern matching?



# Z-value Pattern Matching

To solve pattern matching (given  $P$  and  $T$ ), let  $S = P\$T$

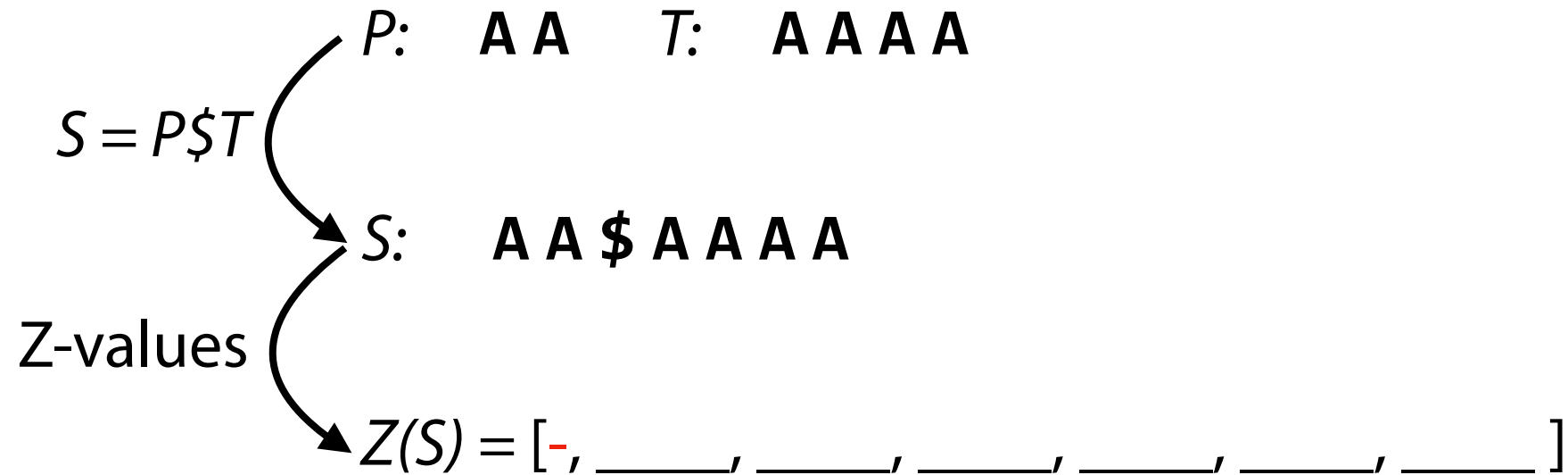
$\$$  = 'terminal character', outside alphabet



# Z-value Pattern Matching

To solve pattern matching (given  $P$  and  $T$ ), let  $S = P\$T$

$\$$  = 'terminal character', outside alphabet



# Z-value Pattern Matching

To solve pattern matching (given  $P$  and  $T$ ), let  $S = P\$T$

$\$$  = 'terminal character', outside alphabet

$P$ : A A     $T$ : A A A A

0 1 2 3 4 5 6  
 $S$ : A A \$ A A A A  
      0 1 2 3

$Z(S) = [-, 1, 0, 2, 2, 2, 1]$

What  $Z_i$  values are matches?

What are the matching indices in  $T$ ?



# Z-value Pattern Matching



$P:$  TT     $T:$  CTTA

## Z-value search pseudo-code

$S:$

1. Concatenate ( $S=P\$T$ )

$Z(S):$

2. Calculate Z-values for  $S$

3. For  $i < 0$ , match if  $Z_i =$  \_\_\_\_\_

Match is **not** at  $i$ , but instead at

\_\_\_\_\_

# Assignment 2: a\_zval

Learning Objective:

Construct a Z-value calculator and measure its efficiency

Demonstrate use of Z-values in pattern matching

Consider: Our goal is  $\theta(|P| + |T|)$ . Does Z-value search match this?

# End-of-class brainstorm

What information does a single Z-value tell us?

If I know  $Z_{i-1}(S)$ , can I use that information to help me compute  $Z_i(S)$ ?



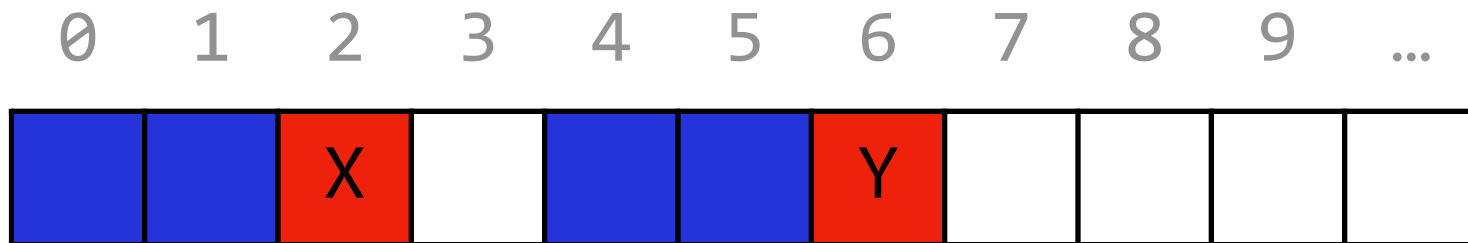
# The Z-value (Take 2)

Given a string  $S$ ,  $Z_i(S)$  is the length of the longest substring in  $S$ , starting at position  $i$ , that matches a prefix of  $S$ .

What information does this give us?

$S$ : **XYXYXABCD**

$$Z_4 = 2$$

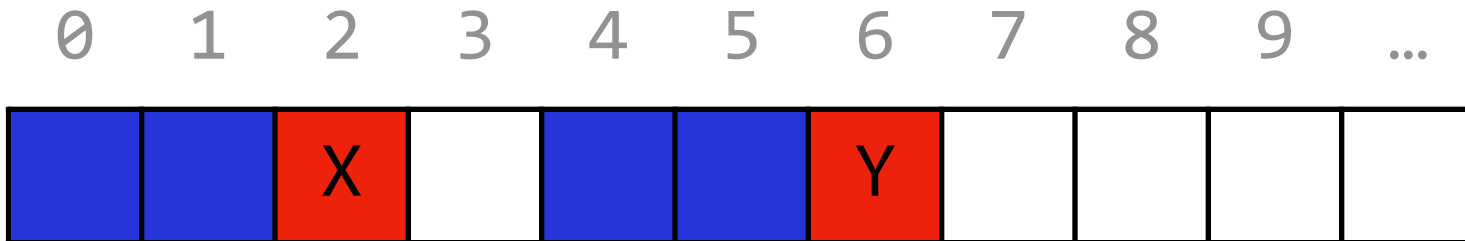


# The Z-value (Take 2)

Given a string  $S$ ,  $Z_i(S)$  is the length of the longest substring in  $S$ , starting at position  $i$ , that matches a prefix of  $S$ .

What information does this give us?

$\begin{matrix} 0 & 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 & 9 \\ S: & \boxed{T} & \boxed{T} & C & G & \boxed{T} & \boxed{T} & A & G & C & G \end{matrix} \quad Z_4 = 2$



# The Z-Algorithm

Assume we've computed  $Z_1, \dots, Z_{i-1}$  and need to calculate  $Z_i$

**Case 1:** We know nothing about the characters at  $S[i]$

$Z_1 = ?$

	0	1	2	3	4	5	6	7
	A	A	A	A	B	B	B	B
	A	A	A	A	B	B	B	B

**Case 2:** We know something about the characters at  $S[i]$

$Z_2 = ?$

	0	1	2	3	4	5	6	7
	A	A	A	A	B	B	B	B
	A	A	A	A	B	B	B	B

# The Z-Algorithm

$$Z_1 = 3$$

$$Z_2 = ?$$

0	1	2	3	4	5	6	7
A	A	A	A	B	B	B	B
A	A	A	A	B	B	B	B

We track our current knowledge of  $S$  using three values:  $i, r, l$

$i$ , the current index position being calculated

$r$ , the index of the rightmost character which has ever been matched

$l$ , the index of Z-value which  $r$  belongs too



# The Z-Algorithm

Start

End

$i$ , the current index =

$r$ , the furthest match char =

$l$ , the furthest reaching Z-value =

-	-	-	-	-	-	-	-
0	1	2	3	4	5	6	7
A	A	B	B	A	A	B	A
A	A	B	B	A	A	B	A

# The Z-Algorithm

Start

End

$i$ , the current index =

$r$ , the furthest match char =

$l$ , the furthest reaching Z-value =

-	1						
0	1	2	3	4	5	6	7
A	A	B	B	A	A	B	A
A	A	B	B	A	A	B	A

# The Z-Algorithm

Start

End

$i$ , the current index =

$r$ , the furthest match char =

$l$ , the furthest reaching Z-value =

-	1	0	0				
0	1	2	3	4	5	6	7
A	A	B	B	A	A	B	A
A	A	B	B	A	A	B	A

# The Z-Algorithm

Start

End

$i$ , the current index =

$r$ , the furthest match char =

$l$ , the furthest reaching Z-value =

-	1	0	0				
0	1	2	3	4	5	6	7
A	A	B	B	A	A	B	A
A	A	B	B	A	A	B	A

# The Z-Algorithm

Start

End

$i$ , the current index =

$r$ , the furthest match char =

$l$ , the furthest reaching Z-value =

-	1	0	0	3			
0	1	2	3	4	5	6	7
A	A	B	B	A	A	B	A
A	A	B	B	A	A	B	A

# The Z-Algorithm

Start

End

$i$ , the current index =

$r$ , the furthest match char =

$l$ , the furthest reaching Z-value =

-	1	0	0	3	1	0	—
0	1	2	3	4	5	6	7
A	A	B	B	A	A	B	A
A	A	B	B	A	A	B	A

# The Z-Algorithm

Start

End

$i$ , the current index =

$r$ , the furthest match char =

$l$ , the furthest reaching Z-value =

-	1	0	0	3	1	0	1
0	1	2	3	4	5	6	7
A	A	B	B	A	A	B	A
A	A	B	B	A	A	B	A

# The Z-Algorithm



**Intuition:** We can use the previous  $Z_1, \dots, Z_i$  to compute  $Z_{i+1}$ !

Track 'what we know' using three integers:  $i, r, l$

Next week: Review how integers are updated to define specific cases.