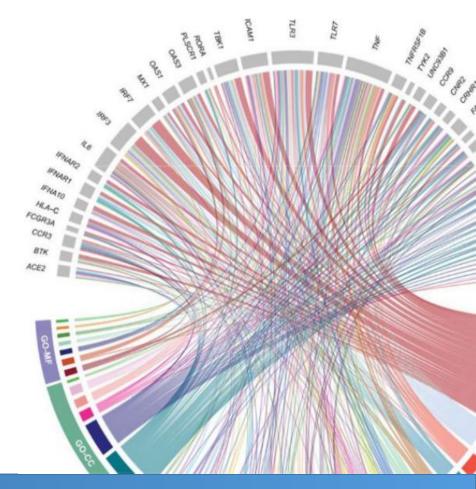
Tècniques i Eines Bioinformàtiques

Boyer-Moore Algorithm

Santiago Marco-Sola (santiago.marco@upc.edu)

Màster en Enginyeria Informàtica, UPC
Departament of Computer Science
Facultat d'Informàtica de Barcelona (FIB), UPC





Acknowledgements

Many pictures and materials are taken from **Ben Langmead's course**.

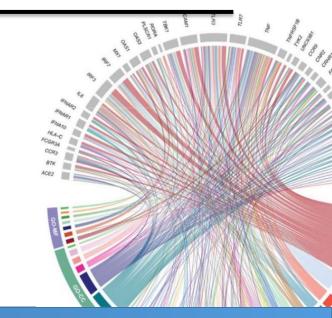
Course heavily inspired in:

- **Genome-Scale Algorithm Design**. Veli Mäkinen, Djamal Belazzougui, Fabio Cunial, Alexandru I. Tomescu. Cambridge University Press.
- Algorithms on Strings, Trees, and Sequences. Dan Gusfield.
 Cambridge University Press.
- An Introduction to Bioinformatics Algorithms. Neil C. Jones, Pavel A. Pevzner. MIT Press.



1

Boyer-Moore Algorithm



Can we improve on the naïve algorithm?

u doesn't occur in P, so skip next two alignments

```
P: word

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

word

word skip!

word skip!

word
```



Boyer-Moore

Learn from character comparisons to skip pointless alignments

1. When we hit a mismatch, move *P* along until the mismatch becomes a match

"Bad character rule"

2. When we move *P* along, make sure characters that matched in the last alignment also match in the next alignment

"Good suffix rule"

3. Try alignments in one direction, but do character comparisons in *opposite* direction

"Longer skips"

P: word

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



Boyer-Moore: Bad character rule

- Upon mismatch, skip alignments until:
 - (a) mismatch becomes a match
 - (b) P moves past mismatched character.
 - (c) If there was no mismatch, don't skip

```
T: GCTTCTGCTACCTTTTGCGCGCGCGCGAA
Step 1:
                                         Case (a)
      T: GCTTCTGCTACCTTTTGCGCGCGCGCGAA
Step 2:
                                         Case (b)
      T: GCTTCTGCTACCTTTTGCGCGCGCGCGAA
Step 3:
                                         Case (c)
      T: GCTTCTGCTACCTTTTGCGCGCGCGCGAA
Step 4:
                     CCTTTTGC
```



Boyer-Moore: Bad character rule

```
Step 1: T: GCTTCTGCTACCTTTTGCGCGCGCGCGCGAA

P: CCTTTTGC

Step 2: T: GCTTCTGCTACCTTTTGCGCGCGCGCGCGAA

Step 3: T: GCTTCTGCTACCTTTTGCGCGCGCGCGCGAA

P: CCTTTTGC

↑ ↑ ↑ ↑ ↑ ↑ ↑ ↑
```

Up to step 3, we skipped 8 alignments

5 characters in Twere never looked at

Boyer-Moore: Good suffix rule

Let t = substring matched by inner loop; skip until (a) there are no mismatches between P and t or (b) P moves past t

```
Step 1: T: CGTGCCTACTTACTTACTTACGCGAA

P: CTTACTTAC

Step 2: T: CGTGCCTACTTACTTACTTACGCGAA

P: CTTACTTAC

Step 3: T: CGTGCCTACTTACTTACTTACGCGAA

CTTACTTAC
```



Boyer-Moore: Good suffix rule

Let t = substring matched by inner loop; skip until (a) there are no mismatches between P and t or (b) P moves past t

```
Step 1: 
T: CGTGCCTACTTACTTACTTACGCGAA

P: CTTACTTAC

toccurs in its entirety to the left within P

Step 2: 
T: CGTGCCTACTTAC

prefix of P matches a suffix of t

Step 3: 
P: CTTACTTAC

TTACTTACTTACTTACGCGAA

CTTACTTACTTACTTACTTACGCGAA

CTTACTTAC
```

Case (a) has two subcases according to whether t occurs in its entirety to the left within P (as in step 1), or a prefix of P matches a suffix of t (as in step 2)



Boyer-Moore: Putting it together

How to *combine* bad character and good suffix rules?

bad char says skip 2, good suffix says skip 7

Take the maximum! (7)

Boyer-Moore: Putting it together

Use bad character or good suffix rule, whichever skips more

```
Step 1: T: GTTATAGCTGATCGCGGCGTAGCGGCGAA

P: GTAGCGGCG

bc: 6, gs: 0 bad character

Step 2: T: GTTATAGCTGATCGCGGCGTAGCGGCGAA

P: GTAGCGGCG

bc: 0, gs: 2 good suffix

Step 3: T: GTTATAGCTGATCGCGGCGTAGCGGCGAA

P: GTAGCGGCG

Step 4: T: GTTATAGCTGATCGCGGCGTAGCGGCGAA

GTAGCGGCGAA

GTAGCGGCGAA

GTAGCGGCG
```



Boyer-Moore: Comparisons Skipped (characters ignored)

11 characters of *T* we ignored

```
T: GTTATAGCTGATCGCGGCGTAGCGGCGAA
Step 1:
      P: GTAGCGGCG
      T: GTTATAGCTGATCGCGGCGTAGCGGCGAA
Step 2:
                GTAGCGGCG
      T: GTTATAGCTGATCGCGGCGTAGCGGCGAA
Step 3:
                    GTAGCGGCG
       GTTATAGCTGATCGCGGCGTAGCGGCGAA
Step 4:
                             GTAGCGGCG
      P:
          Skipped 15 alignments
```



Boyer-Moore: Good suffix rule

We learned the *weak* good suffix rule; there is also a strong good suffix rule

```
7: CTTGCCTACTTACTACT

P: CTTACTTAC

Weak: CTTACTTAC

Strong: CTTACTTAC

guaranteed

mismatch!
```

Strong good suffix rule skips more than weak, at no additional penalty

Strong rule is needed for proof of Boyer-Moore's O(n + m) worst-case time. Gusfield discusses proof(s) in first several sections of ch. 3



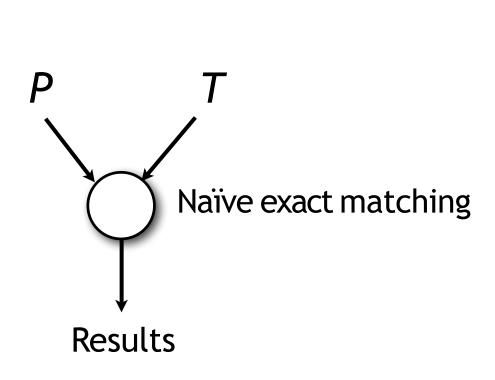
Boyer-Moore implementation

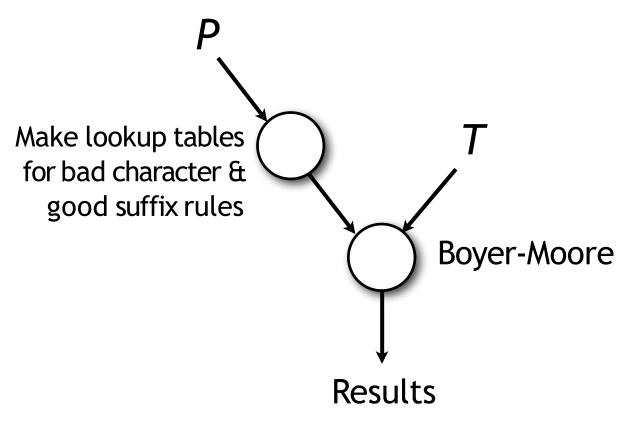
```
Boyer-Moore Algorithm
def boyer_moore(p,p_bm,t):
  # Do Boyer-Moore matching
  i = 0
  occurrences = []
  while i < len(t) - len(p) + 1: # Left to right
    shift = 1
   mismatched = False
   for j in range(len(p)-1,-1,-1): # Right to left
      if p[i] != t[i+i]:
        skip bc = p bm.bad character rule(j,t[i+j])
        skip gs = p bm.good suffix rule(j)
        shift = max(shift,skip bc,skip gs)
        mismatched = True
        break
    if not mismatched:
      occurrences.append(i)
      skip gs = p bm.match skip()
      shift = max(shift,skip_gs)
    i += shift
    return occurrences
```

Full Code:

http://j.mp/CG_BoyerMoore

Preprocessing. Naïve algorithm vs. Boyer-Moore





Boyer-Moore: Preprocessing

- Pre-calculate skips for all possible mismatch scenarios!
- This can be constructed efficiently. See Gusfield 2.2.2.
- As with bad character rule, good suffix rule skips can be precalculated efficiently. See Gusfield 2.2.4 and 2.2.5.
- For both tables, the calculations only consider P. No knowledge of T is required.



-		
- 1	ľ	1
	L	ı
•		

		Т	С	G	С
	Α	0	1	2	3
	С	0	•	0	-
•	G	0	1	•	0
		-	0	1	2



Boyer-Moore: Best case

What's the best case?

- P: bbbb

Every alignment yields immediate mismatch and bad character rule skips *n* alignments

How many character comparisons? floor(m / n)

Boyer-Moore: Worst case

Boyer-Moore, with refinements in Gusfield, is O(n + m) time

Given n < m, can simplify to O(m)

Is this better than naïve?

For naïve, worst-case # char comparisons is n(m - n + 1)

Boyer-Moore: O(m), naïve: O(nm)

Reminder: |P| = n |T| = m

Naïve vs Boyer-Moore

• As m and n grow, the number characters comparisons grows with...

	Naïve matching	Boyer-Moore
Worst case	m∙n	m
Best case	m	m /n

Performance comparison

Simple Python implementations of naïve and Boyer-Moore:

	Naïve matching		Boyer-Moore		
	#character comparisons	wall clock time	#character comparisons	wall clock time	
P: "tomorrow" T: Shakespeare's complete works	5,906,125	2.90 s	785,855	1.54 s	17 matches <i>T</i> = 5.59 M
P: 50 nt string from Alu repeat*	307,013,905	137 s	32,495,111	55 s	336 matches T = 249 M
T : Human reference (hg19) chromosome 1					



^{*} GCGCGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCCGAGGCGGG

Preprocessing: Boyer-Moore

Preprocessing: trade one-time cost for reduced work overall via reuse

Boyer-Moore preprocesses *P* into lookup tables that are *reused*

reused for each alignment of P to T₁

If you later give me T_2 , I reuse the tables to match P to T_2

If you later give me T_3 , I reuse the tables to match P to T_3

•••

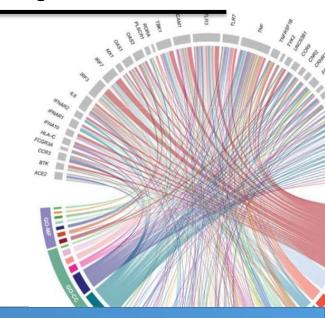
Cost of preprocessing is amortized over alignments & texts





3

Exercises (Hands-on)



Exercises

1. Implement the Bad Character Rule (Easy)

• Implement the bad character rule, one of the optimizations of Boyer-Moore. Create a function that precomputes the last occurrence of each character in P. Use this table to determine the shift when a mismatch occurs.

2. Combine Naïve Algorithm and Bad Character Rule (Intermediate)

• Find all occurrences of a short pattern (read) within a longer reference genome sequence. Modify the naïve string matching function to incorporate the bad character rule. Test it with a small reference genome.

3. Implement the Good Suffix Rule (Advanced)

• Find all occurrences of a short pattern (read) within a longer reference genome sequence. Implement a simple exact matching algorithm.

4. Full Boyer-Moore Implementation (Expert)

 Implement the complete Boyer-Moore algorithm using both the bad character and good suffix rules. Combine the bad character and good suffix heuristics to achieve efficient string matching. Compare performance against the naïve algorithm.



Helpers

Download chr1.fa

> wget http://hgdownload.soe.ucsc.edu/goldenPath/hg38/chromosomes/chr1.fa.gz

Parsing and Reading Chr1.fa

```
import argparse
def read_fasta(file_path):
 sequence = []
 with open(file_path, 'r') as file:
   for line in file:
     if not line.startswith('>'):
       sequence.append(line.strip()) # Remove newline and concatenate
 return ''.join(sequence)
if name == " main ":
 parser = argparse.ArgumentParser(
   description="Read a FASTA file and store the sequence in a string.")
 parser.add argument("i", help="Path to the FASTA file")
 args = parser.parse args()
 sequence = read fasta(args.fasta file)
```



Exercise 1: Implement the Bad Character Rule

1. Implement the Bad Character Rule (Easy)

• Implement the bad character rule, one of the optimizations of Boyer-Moore. Create a function that precomputes the last occurrence of each character in P. Use this table to determine the shift when a mismatch occurs.

Bad Character Rule import string def bad_character_table(pattern): Precompute the last occurrence table for each character in the pattern. bad char = {} for i in range(len(pattern)): bad char[pattern[i]] = i # Store the last occurrence of each character return bad char def bad character shift(pattern, mismatch char, mismatch index): Determine the shift when a mismatch occurs. bad char = bad character table(pattern) last occurrence = bad char.get(mismatch char, -1) # Default to -1 if character not in pattern shift = max(1, mismatch index - last occurrence) # Shift to align the next best possible match return shift



Exercise 1: Implement the Bad Character Rule

1. Implement the Bad Character Rule (Easy)

• Implement the bad character rule, one of the optimizations of Boyer-Moore. Create a function that precomputes the last occurrence of each character in P. Use this table to determine the shift when a mismatch occurs.

Bad Character Rule

```
# Example usage
pattern = "GCTTAC"
text = "AAGCTTGCCTTACGCTTAC"

# Generate the bad character table
bad_char_table = bad_character_table(pattern)
print("Bad Character Table:", bad_char_table)

# Simulate a mismatch at index 4 with character 'G'
mismatch_char = "G"
mismatch_index = 4
shift = bad_character_shift(pattern, mismatch_char, mismatch_index)
print(f"Mismatch at index {mismatch_index} with char '{mismatch_char}', shift by {shift} positions.")
```

Exercise 2: Combine Naïve Algorithm and Bad Character Rule

Naïve Algorithm and Bad Character Rule **Testing** def boyer moore bad character(text, pattern): Boyer-Moore exact string matching using the Bad Character Rule.Finds all occurrences of `pattern` in `text`. m, n = len(pattern), len(text) if m > n: return [] # Pattern is longer than text, no matches possible bad char = bad character table(pattern) # Precompute bad character table occurrences = [] i = 0 # Start index in text while i <= n - m: j = m - 1 # Start checking from the end of the pattern # Compare pattern with text from right to left while j >= 0 and pattern[j] == text[i + j]: j -= 1 if j < 0: # Pattern found at index i occurrences.append(i) # Shift pattern i += (m - bad char.get(text[i + m], -1)) if i + m < n else 1else: # Mismatch occurred, apply the Bad Character Rule shift = max(1, j - bad_char.get(text[i + j], -1)) i += shift # Move pattern to the right by the shift return occurrences

```
# Example usage
text = "AGCTTAGCTAAGCTAGCTAGCTA"

pattern = "AGCTA"

matches = boyer_moore_bad_character(text, pattern)
print(f"Pattern found at positions: {matches}")
```

