Burrows-Wheeler Transform

Methods in Bioinformatics (TI)



Luca Denti

Strings

Alphabet Σ

String S

String: AAGTGCTCAAAGCTAAGCTCCAT

Prefix:

Suffix:

Substring:

Strings

• Alphabet Σ : set of *characters* (e.g., Σ ={A,C,G,T})

• **String S**: sequence of n=|S| characters drawn from Σ , i.e., S[i] $\in \Sigma$ for $0 \le i < n$

String: AAGTGCTCAAAGCTAAGCTCCAT

Prefix: AAGTGC

Suffix: CAT

Substring: AAAGC

String Ordering

Lexicographic/alphabetical order

animal < house < ta < tac < zoo

When no character breaks the tie (e.g., one string is prefix of the other), shorter comes first.

String rotation refers to the process of moving characters in a string from one end to the other while maintaining their order.

TACTAC

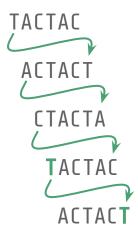
String rotation refers to the process of moving characters in a string from one end to the other while maintaining their order.

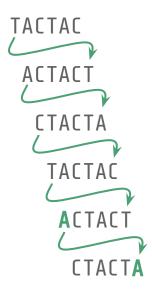
TACTAC ACTACT

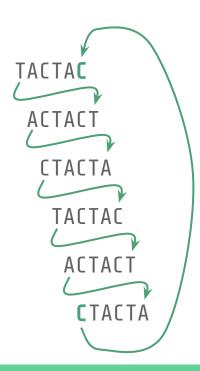
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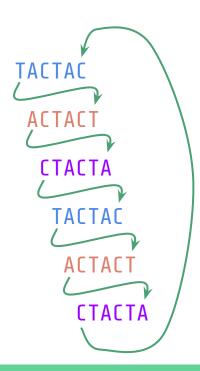
ACTACT CTACTA











Special character \$

- 1. Define a new symbol **\$**:
 - \$∉Σ
 - \$<c ∀c∈Σ
 </p>
- 2. Append \$ to the string

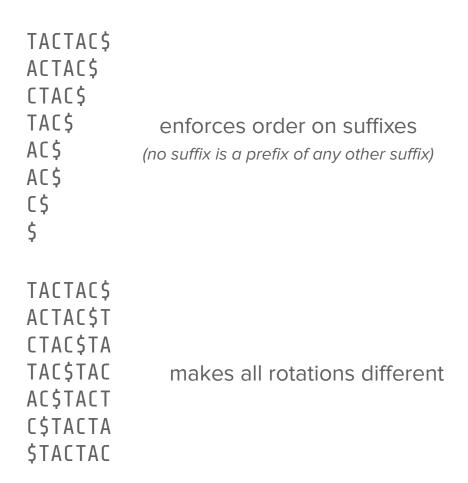
Special character \$

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 - \$∉Σ
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Special character \$

- 1. Define a new symbol \$:

 - $\$<c \forall c \in \Sigma$
- 2. Append \$ to the string



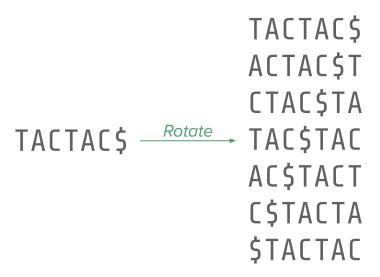
Reversible permutation of the characters of a string, introduced for **compression**

Reversible permutation of the characters of a string, introduced for **compression**(i) (ii) (iii)

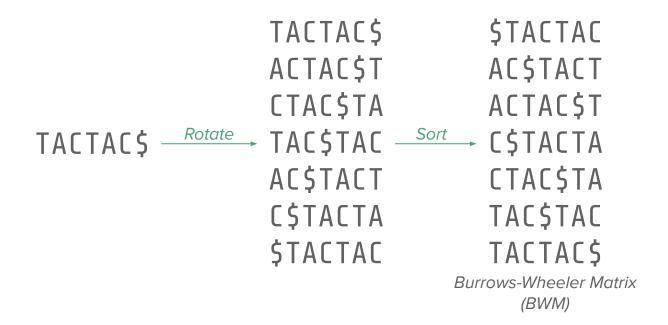
Reversible permutation of the characters of a string, introduced for **compression**(i) (iii) (iii)

TACTAC\$

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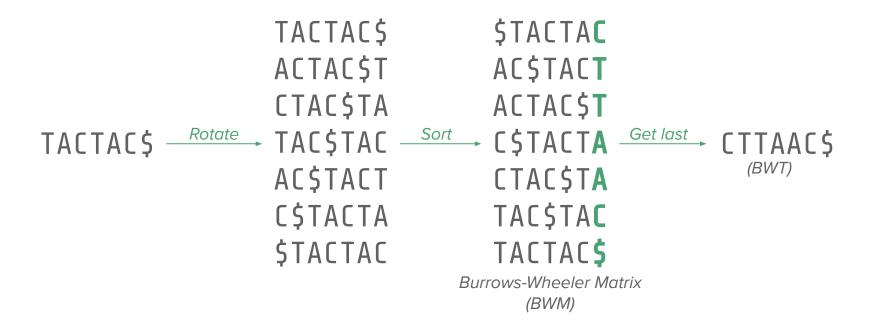


Reversible permutation of the characters of a string, introduced for **compression**(i) (iii) (iii)



Burrows M, Wheeler DJ: A block sorting lossless data compression algorithm. Digital Equipment Corporation, Palo Alto, CA 1994, Technical Report 124; 1994

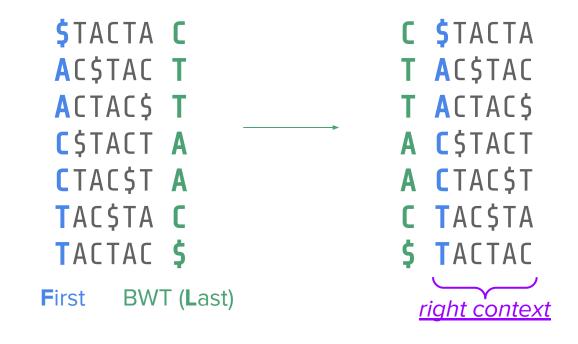
Reversible permutation of the characters of a string, introduced for **compression**(i) (iii) (iii)



Burrows M, Wheeler DJ: A block sorting lossless data compression algorithm. Digital Equipment Corporation, Palo Alto, CA 1994, Technical Report 124; 1994

Permutation

BWT permutes characters according to their right contexts



Compression

BWT <u>facilitates</u> compression <u>(it does not compress the input string)</u>

- it tends to cluster identical characters together
- it combines repeated patterns into larger contiguous blocks
- it makes the string easier to compress (e.g., run-length encoding)

Compression

BWT <u>facilitates</u> compression <u>(it does not compress the input string)</u> - **bzip**

- it tends to cluster identical characters together
- it combines repeated patterns into larger contiguous blocks
- it makes the string easier to compress (e.g., run-length encoding)



> 14,A (9 bytes vs 14 bytes)*

*Very rough approximation, implementation-dependent, no encoding (2bit/packed)

BWT in Bioinformatics

- especially convenient for short reads
 - o millions of string searches in a long string
- query complexity depends on read size (not on genome size)
- "construction" is a 1 time expense

Ultrafast and memory-efficient **alignment** of short DNA sequences to the human genome

B Langmead, C Trapnell, M Pop, SL Salzberg - Genome biology, 2009 - Springer

... For the human genome, Burrows-Wheeler indexing allows **Bowtie** to **align** more than 25 million reads per CPU hour with a memory footprint of approximately 1.3 gigabytes. ...

☆ Salva ワワ Cita Citato da 25169 Articoli correlati Tutte e 29 le versioni

Fast and accurate short read alignment with Burrows-Wheeler transform

H Li, R Durbin - bioinformatics, 2009 - academic.oup.com

..., and present the algorithm for inexact matching which is implemented in **BWA**. We evaluate the performance of **BWA** on simulated data by comparing the **BWA alignment** with the true ...

☆ Salva 兒 Cita Citato da 52831 Articoli correlati Tutte e 29 le versioni

STAR: ultrafast universal RNA-seq aligner

A Dobin, CA Davis, F Schlesinger, J Drenkow... - ..., 2013 - academic.oup.com

... **Alignment** to a Reference (**STAR**) software based on a previously undescribed RNA-seq **alignment** ... **STAR** outperforms other aligners by a factor of >50 in mapping speed, aligning to the ...

☆ Salva ワワ Cita Citato da 50265 Articoli correlati Tutte e 21 le versioni

<u>Reversible</u>

$$S \rightarrow BWT(S) \rightarrow S$$



LF-Mapping (Last-to-First)

Property of BWT that allows to reconstruct the original string from the BWT, starting from its end and going backward

BWT(BANANA)

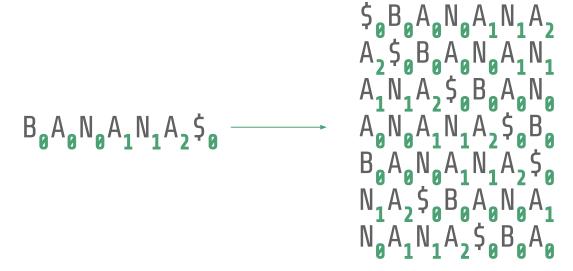
BWT(BANANA)

ANNB\$AA

Let's give each character, its **rank** (number of occurrences up to its position)

$$B_0A_0N_0A_1N_1A_2$$

Let's give each character, its rank (number of occurrences up to its position)



Let's give each character, its **rank** (number of occurrences up to its position)

\$0	A ₂
A ₂	N_1^2
A ₁	N
A	B
B	\$0
N ₁	A ₁
N ₁	A
F	L
	_

We do not need the entire matrix

Let's look at F:

\$ 0	A2
A ₂	N_1^2
A ₁	N
A	B
B	\$0
N ₁	A
N _e	A
F	L

Let's look at F:

- sorted column
- predictable column (as long as we know how many times each character occur)

0		
-		
1		
^ ^		
100		
U		
2		
\		
1 _		
• ¬		
1		
l .		
1		
1 -		
-1		
1		
_		
1		
0		
4		
' O		
170		
9		
,		
١		
<i>,</i>		
•		
1		
1		
0		
_		
⊢		
'		

Let's look at F:

- sorted column
- predictable column (as long as we know how many times each character occur)

Let's look at F and L:

0		
2		
1		
0		
0		
1		
0		
=		



Let's look at F:

- sorted column
- predictable column (as long as we know how many times each character occur)

Let's look at F and L:

As occur in the same order

A

F

LF-Mapping

Let's look at F:

- sorted column
- predictable column (as long as we know how many times each character occur)

Let's look at F and L:

- As occur in the same order
- Same for Bs

3

: L

LF-Mapping

Let's look at F:

- sorted column
- predictable column (as long as we know how many times each character occur)

Let's look at F and L:

- As occur in the same order
- Same for Bs
- Same for Ns

N₁ N₀ N₁

L

LF-Mapping

More generally,

the ith occurrence of a character in L and the ith occurrence of a character in F, correspond to the same occurrence in the original string (i.e., they have the same rank)

1	A
	N,
	N
1	
	B ₀ \$ ₀ A ₁
	A
	A
	L

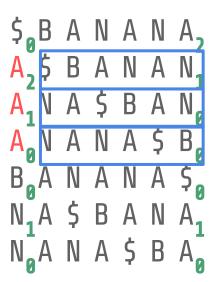
\$ ₀ B	A	N	A	N	Α,
A ₂ \$	В	A	N	A	N ₁
A_1^{N}	A	\$	В	A	N
AN	A	\mathbb{N}	A	\$	B
BOA	N	A	N	A	\$
N_1A	\$	В	A	N	A ₁
N ₀ A	N	A	\$	В	A

Why are these As in this relative order?

Why are these As in this relative order?

They are sorted by their right context

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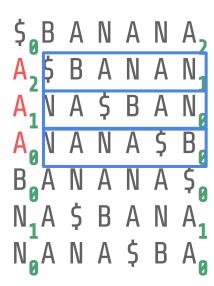


They are sorted by their right context

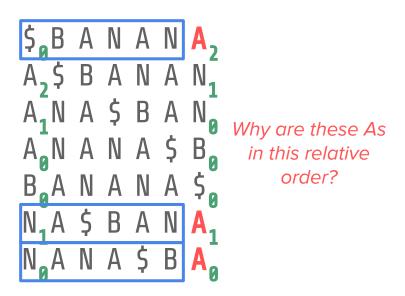


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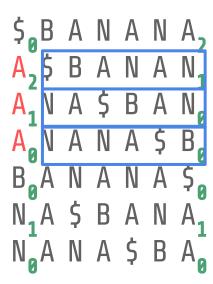


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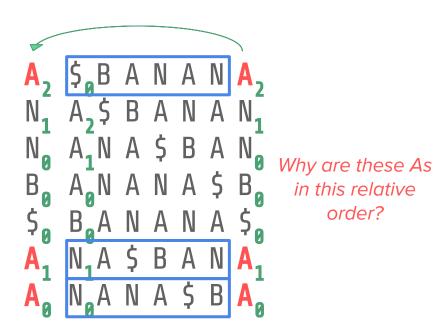


They are sorted by their left context

Why are these As in this relative order?

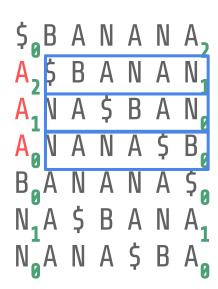


They are sorted by their right context

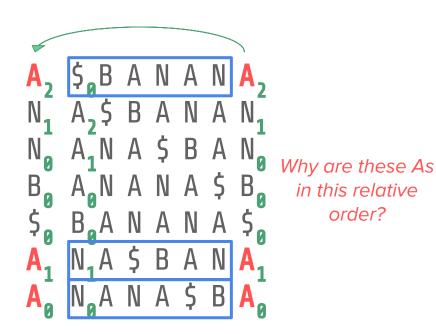


They are sorted by their left context, that by construction (rotations) it's their right context

Why are these As in this relative order?



They are sorted by their right context



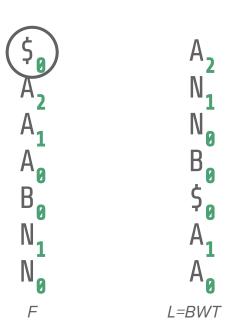
They are sorted by their left context, that by construction (rotations) it's their right context

Both columns are sorted following the same principle, therefore are in the same order

These are just arrays

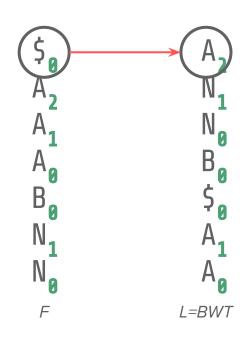
\$0	A ₂
A ₂	N_1^2
A ₁	N
A	B_{0}^{o}
B	\$0
N_{1}	A
N ₀	A
F	L=BW7

1. Start from first row (\$ in F, by construction)

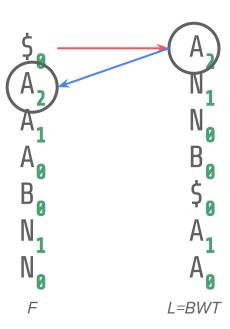


\$0

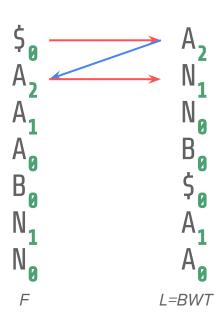
- 1. Start from first row (\$ in F, by construction)
- 2. Move to L, it contains the character preceding \$ (by construction)



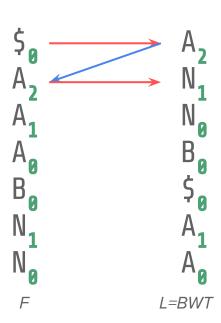
- 1. Start from first row (\$ in F, by construction)
- 2. Move to L, it contains the character preceding \$ (by construction)
- 3. **Jump** to F using LF-Mapping (same rank)



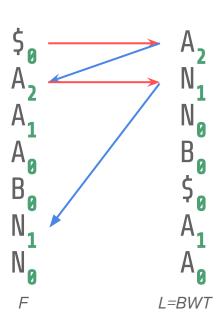
- 1. Start from first row (\$ in F, by construction)
- 2. Move to L, it contains the character preceding \$ (by construction)
- 3. **Jump** to F using LF-Mapping (same rank)
- 4. **Move** to L, it contains the preceding character



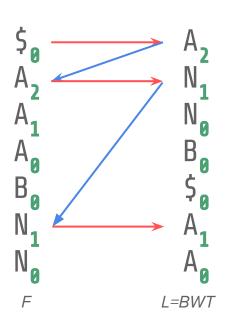
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- 3. **Jump** to F using LF-Mapping (same rank)
- 4. **Move** to L, it contains the preceding character
- 5. Repeat 3-4 until reaching \$ in L



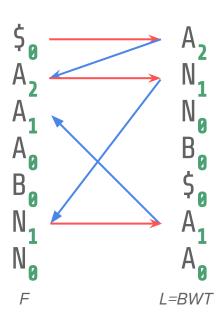
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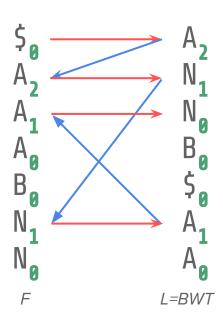
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- 4. **Move** to L, it contains the preceding character
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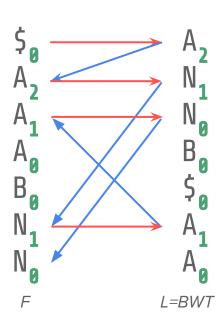
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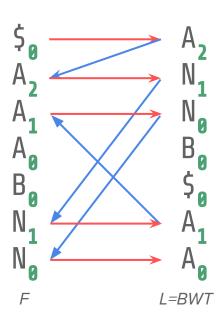
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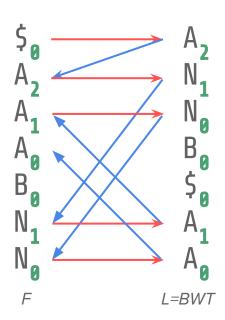


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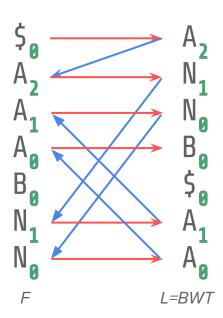
$$A_0 N_0 A_1 N_1 A_2 S_0$$

- 1. Start from first row (\$ in F, by construction)
- 2. Move to L, it contains the character preceding \$ (by construction)
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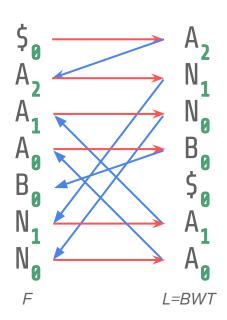
$$A_0 N_0 A_1 N_1 A_2 S_0$$

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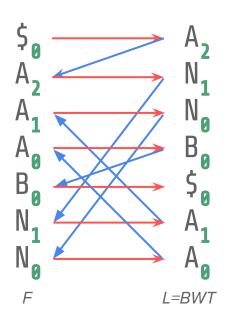
$$B_0A_0N_0A_1N_1A_2$$
\$

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- 3. **Jump** to F using LF-Mapping (same rank)
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$$B_0A_0N_0A_1N_1A_2$$
\$

Given a text T and a pattern P, find P in T

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Three queries:

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• exist: does P occur in T? Yes/no

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Given a text T and a pattern P, find P in T

Three queries:

• exist: does P occur in T? Yes/no

• **count:** how many times does P occur in T? 3

• **locate:** where does P occur in T? Positions 2 and 5

Naive solution:

$$|T| = n$$

 $|P| = m$

Advanced algorithms:

Index-based algorithms (very useful in bioinformatics)

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|T| = n|P| = m

• check for P at every position in T O(n*m)

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Advanced algorithms:

• <u>Knuth-Morris-Pratt</u> O(n + m)

• Boyer-Moore O(n/m) on average, O(n*m) in worst case

• Rabin-Karp O(n + m) on average, $O(n^*m)$ worst case

Index-based algorithms (very useful in bioinformatics)

Naive solution:

|T| = n|P| = m

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Advanced algorithms:

• Knuth-Morris-Pratt O(n + m)

• Boyer-Moore O(n/m) on average, O(n*m) in worst case

• Rabin-Karp O(n + m) on average, $O(n^*m)$ worst case

Index-based algorithms (very useful in bioinformatics)

• FM-Index (BWT-based) O(n) for construction (one time expense), O(m) for matching*

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

```
1 $ A
2 A $ N
3 A N A $ N
4 A N A N A $ B
5 B A N A N A $ A
7 N A N A $ A
6 L=BWT
```

Algorithm is based on:

What are these?

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

1 \$ A 2 A \$ N 3 A N A \$ N 4 A N A N A \$ B 5 B A N A N A \$ 6 N A \$ A 7 N A N A \$ A EBW

Algorithm is based on:

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

What are these?
Lexicographically ordered suffixes

2 A\$\$ N

4 A N A N A \$ B

5 B A N A N A \$

6 N A \$

7 N A N A \$

LERW

Algorithm is based on:

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

What are these? Lexicographically ordered suffixes

1	\$						A
2	A	\$					N
3	A	N	A	\$			N
4	A	N	A	N	A	\$	В
5	В	A	N	A	N	A	\$
6	N	A	\$				A
7	N	A	N	A	\$		A
	F					L=L	BWT

B A N A N A \$
0 1 2 3 4 5 6

Algorithm is based on:

• Q-intervals: intervals on the F column referring to string Q

6 1 \$

• **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

		_						
	2	A	\$					N
	3	A	N	A	\$			N
What are these?	4	A	N	A	N	A	\$	В
Lexicographically ordered suffixes	5	В	A	N	A	N	A	\$
Samiles	6	N	A	\$				A
	7	N	Α	N	Α	\$		A
		F					1 =	RW/

B A N A N A \$

9 1 2 3 4 5 6

Algorithm is based on:

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

5 2 A \$ N

What are these?

Lexicographically ordered suffixes

6 N A \$ A N A N A \$ A

7 N A N A \$ A

L=BWT

B A N A N A \$
0 1 2 3 4 5 6

Algorithm is based on:

- Q-intervals: intervals on the F column referring to string Q
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What are these?
Lexicographically ordered suffixes

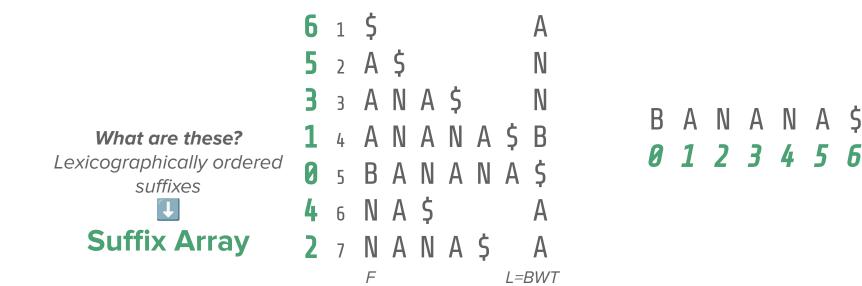
6 N A \$ A

7 N A N A \$ A

L=BWT

B A N A N A \$
0 1 2 3 4 5 6

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)



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- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

	1	\$						A
	2	Α	\$					N
B-interval:	3	A	N	A	\$			N
BAN-interval: A-interval:	4	A	N	A	N	A	\$	В
NA-interval:	5	В	A	N	A	N	A	\$
	6	N	A	\$				Α
	7	N	A	N	A	\$		Α
		F					L=I	BWT

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

	1	\$						A
	2	A	\$					N
B-interval: [5,5]	3	A	N	A	\$			N
BAN-interval: A-interval:	4	A	N	A	N	A	\$	В
NA-interval:	5	В	A	N	A	N	A	\$
	6	N	A	\$				Α
	7	N	A	N	A	\$		Α
		F					L=	BWT

Algorithm is based on:

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

L=BWT

	1	7						A
	2	A	\$					N
B-interval: [5,5]	3	A	N	A	\$			N
BAN-interval: [5,5] A-interval:	4	A	N	A	N	A	\$	В
NA-interval:	5	В	A	N	A	N	A	\$
	6	N	A	\$				A
	7	N	A	N	A	\$		A

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

	2
B-interval: [5	,5]
BAN-interval: [5	
A-interval: [2	[,4]
NA-interval:	5

```
3 A N A $
4 ANANA$B
5 BANANA$
6 N A $
7 NANA$
          L=BWT
```

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

	2	Α	\$			
B-interval: [5,5]	3	A	N	Α	\$	
BAN-interval: [5,5] A-interval: [2,4]	4	A	N	A	N	
NA-interval: [6,7]	5	В	A	N	A	
	6	N	A	\$		

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

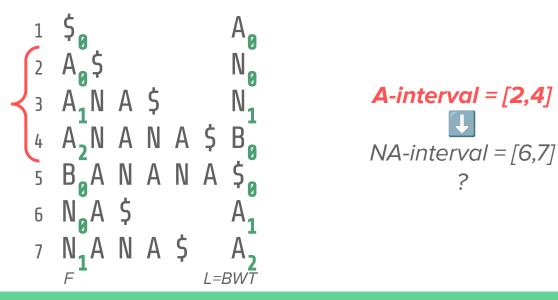
B-interval:	[5,5]
BAN-interval	: [5,5]
A-interval:	[2,4]
NA-interval:	[6,7]

```
3 A N A $
4 A N A N A $ B
5 B A N A N A $
6 N A S
7 NANA$
            L=BWT
```

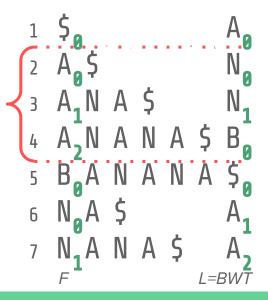
- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)

1	۶ ₀					A
2	A ₀ \$					N
3	A_1N	A	\$			N ₁
	$A_2^{-}N$			A	\$	B
	BA					\$ 0
6	N _a A	_				A ₁
7			A	\$		A ₂ BWT
	F				L=L	BWT

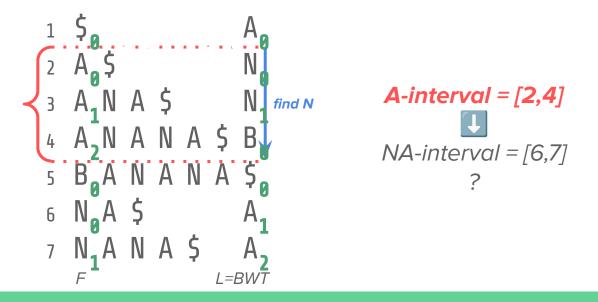
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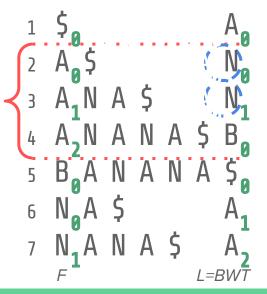


- Q-intervals: intervals on the F column referring to string Q
- LF-mapping: how to obtain cQ-interval from Q-interval (backward extension)



Algorithm is based on:

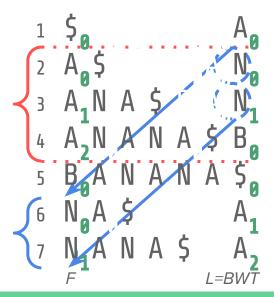
- Q-intervals: intervals on the F column referring to string Q
- LF-mapping: how to obtain cQ-interval from Q-interval (backward extension)



same character in L are not always contiguous

Algorithm is based on:

- Q-intervals: intervals on the F column referring to string Q
- **LF-mapping:** how to obtain cQ-interval from Q-interval (backward extension)



same character in L are not always contiguous but thanks to LF-mapping, they are on F

We can search a pattern P via IPI backward extensions

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We can search a pattern P via |P| backward extensions

A-interval = [2,4]

We can search a pattern P via |P| backward extensions

A-interval = [2,4] \rightarrow NA-interval = [6,7]

We can search a pattern P via |P| backward extensions

A-interval = [2,4] \Rightarrow NA-interval = [6,7]

We can search a pattern P via |P| backward extensions

A-interval = [2,4] \rightarrow NA-interval = [6,7] \rightarrow ANA-interval [3,4]

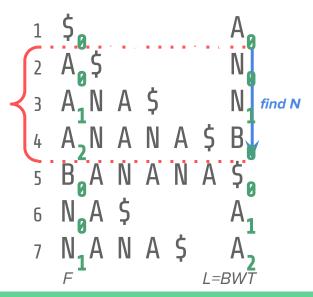
We can search a pattern P via IPI backward extensions

A-interval = [2,4] \Rightarrow NA-interval = [6,7] \Rightarrow ANA-interval [3,4] \Rightarrow BANA-interval [5,5]

O(m) where m is the length of pattern P

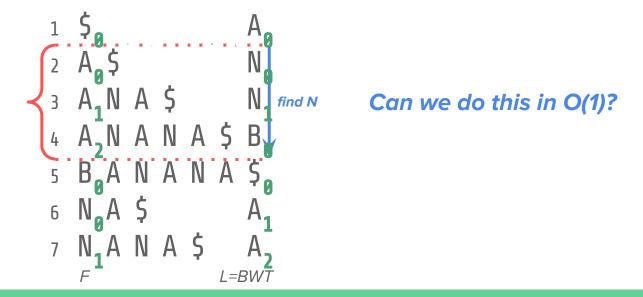
O(m) where m is the length of pattern P

but this is true if we do not need to iterate over each interval to find the character we are interested in!



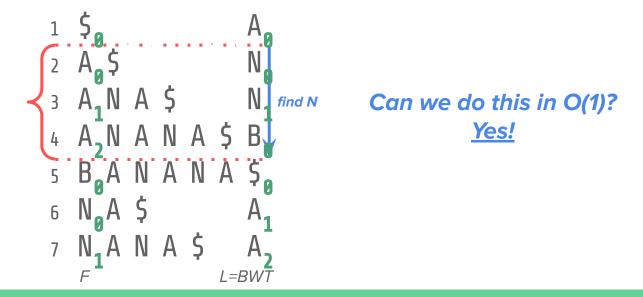
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O(m) where m is the length of pattern P

but this is true if we do not need to iterate over each interval to find the character we are interested in!



FM-Index

- Full-text index combining the BWT with auxiliary data structures
 - efficient indexing
 - efficient querying
 - "store" full input
- Main idea: represent F and L in an efficient and compact way
- Potentially very space-efficient (implementation-dependent)

Efficient backward extension (FM-Index)

Only things we need for backward extensions/search are F and L columns, but we can represent them in a more convenient way

Array C with "cumulative counts" of smaller symbols for each $c \in \{\$\} \cup \Sigma$

```
4 A, N A N A $
```

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Array C with "cumulative counts" of smaller symbols for each $c \in \{\$\} \cup \Sigma$

$$C = [\mathbf{0},]$$
How many symbols we have smaller than \$?

```
4 A N A N A $
```

Only things we need for backward extensions/search are F and L columns, but we can represent them in a more convenient way

Array C with "cumulative counts" of smaller symbols for each $c \in \{\$\} \cup \Sigma$

How many symbols we have smaller than A?

```
4 A N A N A $
```

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Array C with "cumulative counts" of smaller symbols for each $c \in \{\$\} \cup \Sigma$

How many symbols we have smaller than A?

```
4 A N A N A $
```

Only things we need for backward extensions/search are F and L columns, but we can represent them in a more convenient way

Array C with "cumulative counts" of smaller symbols for each $c \in \{\$\} \cup \Sigma$

How many symbols we have smaller than B?

```
4 A N A N A $
```

Only things we need for backward extensions/search are F and L columns, but we can represent them in a more convenient way

Array C with "cumulative counts" of smaller symbols for each $c \in \{\$\} \cup \Sigma$

How many symbols we have smaller than B?

```
4 A N A N A $
```

Only things we need for backward extensions/search are F and L columns, but we can represent them in a more convenient way

Array C with "cumulative counts" of smaller symbols for each $c \in \{\$\} \cup \Sigma$

$$C = [0, 1, 4,]$$

How many symbols we have smaller than N?

```
4 A N A N A $
```

Only things we need for backward extensions/search are F and L columns, but we can represent them in a more convenient way

Array C with "cumulative counts" of smaller symbols for each $c \in \{\$\} \cup \Sigma$

$$C = [0, 1, 4, 5]$$

How many symbols we have smaller than N?

```
4 A N A N A $
```

Only things we need for backward extensions/search are F and L columns, but we can represent them in a more convenient way

Array C with "cumulative counts" of smaller symbols for each $c \in \{\$\} \cup \Sigma$

$$C = [0, 1, 4, 5]$$

1	\$0					A
2	A ₀ \$					N
3	A_1N	A	\$			N,
4	A _N	A	N	A	\$	B
5	BaA	N	A	N	A	\$
6		\$				A
7	N_1A		A	\$		A 8 <i>W</i> 7
	F				L=L	BW7

Rank matrix Occ

Rank Matrix Occ

Occ is a matrix $|\Sigma| \times |T|$ that stores for each position i on BWT(T) and for each character $c \in \Sigma$, the counts the occurrences of c in the first i elements of BWT(T)

	BWT	\$ A	В	N
1	А			
2	N			
3	N			
4	В			
5	\$			
6	А			
7	А			

Rank Matrix Occ

Occ is a matrix $|\Sigma| \times |T|$ that stores for each position i on BWT(T) and for each character $c \in \Sigma$, the counts the occurrences of c in the first i elements of BWT(T)

	BWT	\$	A	В	N
1	А	0	1	0	0
2	N	0	1	0	1
3	N	0	1	0	2
4	В	0	1	1	2
5	\$	1	1	1	2
6	А	1	2	1	2
7	А	1	3	1	2

How to backward extend using C and Occ?

Given Q-interval [i,j] and symbol c, return cQ-interval if it exists, empty interval otherwise

```
def backwardExtend (c, [i, j]):
    i = C[c] + Occ(c, i - 1) + 1
    j = C[c] + Occ(c, j)
    return [i, j]
```

```
def backwardExtend (c, [i, j]):
    i = C[c] + Occ(c, i - 1) + 1
    j = C[c] + Occ(c, j)
    return [i, j]
```

1	٦					A
2	A ₀ \$					N
	A_1N	A	\$			N ₁
4	A ₂ N	A	N	A	\$	B
5	BA	N	A	N	A	\$
6	N _o A					A ₁
7	N_1A		A	\$		A ₂
	F				L=	BWT

		_			
	BWT	\$	A	В	N
1	А	0	1	0	0
2	N	0	1	0	1
3	N	0	1	0	2
4	В	0	1	1	2
5	\$	1	1	1	2
6	А	1	2	1	2
7	А	1	3	1	2
(С	0	1	4	5

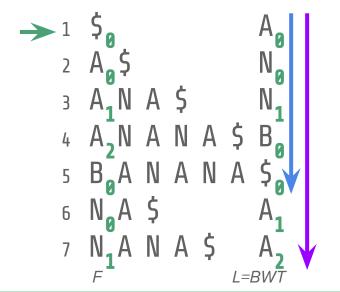
```
def backwardExtend (c, [i, j]):
    i = C[c] + Occ(c, i - 1) + 1
    j = C[c] + Occ(c, j)
    return [i, j]
```

	BWT	\$	Α	В	N
1	А	0	1	0	0
2	N	0	1	0	1
3	N	0	1	0	2
4	В	0	1	1	2
5	\$	1	1	1	2
6	А	1	2	1	2
7	А	1	3	1	2
	С	0	1	4	5

def backwardExtend (c, [i, j]):

$$i = C[c] + Occ(c, i - 1) + 1 = 1 + 1 + 1 = 3$$

 $j = C[c] + Occ(c, j) = 1 + 3 = 4$



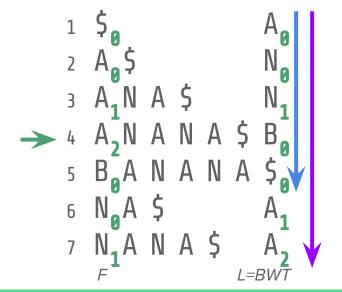
	BWT	\$	Α	В	N
1	А	0	1	0	0
2	N	0	1	0	1
3	N	0	1	0	2
4	В	0	1	1	2
5	\$	1	1	1	2
6	А	1	2	1	2
7	А	1	3	1	2
	С	0	1	4	5

def backwardExtend (c, [i, j]):

$$i = C[c] + Occ(c, i - 1) + 1 = 4 + 1 + 1 = 6$$

 $j = C[c] + Occ(c, j) = 4 + 1 = 5$

return [i, j]



	BWT	\$	Α	В	N
1	А	0	1	0	0
2	N	0	1	0	1
3	N	0	1	0	2
4	В	0	1	1	2
5	\$	1	1	1	2
6	А	1	2	1	2
7	А	1	3	1	2
(С	0	1	4	5

def backwardExtend (c, [i, j]):

$$i = C[c] + Occ(c, i - 1) + 1 = 5 + 1 + 1 = 7$$

 $j = C[c] + Occ(c, j) = 5 + 2 = 7$
return [i, j]

	BWT	\$	Α	В	N
1	А	0	1	0	0
2	N	0	1	0	1
3	N	0	1	0	2
4	В	0	1	1	2
5	\$	1	1	1	2
6	А	1	2	1	2
7	А	1	3	1	2
-	С	0	1	4	5

Backward Search

Given pattern P, find it in T

```
def backwardSearch (P):
    p = len(P)-1
    i,j = C[P[p]], C[P[p]-1] \# assuming order
   while p \ge 0 and i \ge j:
        i,j = backwardExtend(P[p], (i,j))
        p -= 1
   if p >= 0:
        print("P not found")
    else:
        print(f"P found: {j-i+1} occurrences")
```

Not covered here: how to locate occurrences?

Backward Search

Given pattern P, find it in T

```
def backwardSearch (P):
                                                     6 1 $
   p = len(P)-1
                                                     5 2 A S
   i,j = C[P[p]], C[P[p]-1] \# assuming order
   while p \ge 0 and i \ge j:
                                                     3 3 A N A $ N
      i,j = backwardExtend(P[p], (i,j))
                                                     1 4 A N A N A $ B
      p -= 1
                                                     0 5 B A N A N A $
   if p >= 0:
                                                     4 6 N A $
      print("P not found")
                                                     2 7 N A N A S A
   else:
      print(f"P found: {j-i+1} occurrences")
```

Not covered here: how to locate occurrences? Use Suffix Array (although quite expensive, O(nlog(n))

Pattern matching with the FM-Index - Complexity

Query time: O(1) for backward extension, O(m) for backward search

Space: O(n*m)

...but space can be reduced using advanced data structures based on bit vectors:

- wavelet tree
- rope

Not covered here: how to construct BWT/FM-Index

- $O(n^2 \log(n))$
- Vast literature on O(n) approaches
- Start from Suffix Array, O(n) with larger constants

+ what about approximate matches?

	1	\$	С
<u>Bigger example</u>	2	A	G
	3	Α	C
	4	Α	C
	5	C	Τ
	6	C	G
	7	C	C
	8	C	Α
	9		G
	10		G
	11	C	G
	12	C	\$
	13	G	Α
	14	G	C
	15	G	C
	16	G	C
	17	G	C
	18	G	Α
	19	Τ	Τ
	20	Τ	Τ
	21	Τ	G

Bigger example

- 1 \$CGCGCGCGCAGACCAGTTTC
- 2 ACCAGTTTC\$CGCGCGCGCAG
- 3 AGACCAGTTTC\$CGCGCGCGC
- 4 AGTTTC\$CGCGCGCGCAGACC
- 5 C\$CGCGCGCGCAGACCAGTTT
- 6 CAGACCAGTTTC\$CGCGCGCG
- 7 CAGTTTC\$CGCGCGCGCAGAC 8 CCAGTTTC\$CGCGCGCGCAGA
- 9 CGCAGACCAGTTTC\$CGCGCG
- 10 CGCGCAGACCAGTTTC\$CGCG
- 11 CGCGCGCAGACCAGTTTC\$CG
- 12 CGCGCGCGCAGACCAGTTTC\$
- 13 GACCAGTTTC\$CGCGCGCGCA
- 14 GCAGACCAGTTTC\$CGCGCGC
- 15 GCGCAGACCAGTTTC\$CGCGC
- 16 GCGCGCAGACCAGTTTC\$CGC
- 17 GCGCGCGCAGACCAGTTTC\$C 18 GTTTC\$CGCGCGCGCAGACCA
- 19 TC\$CGCGCGCAGACCAGTT
- 20 TTC\$CGCGCGCGCAGACCAGT
- 21 TTTC\$CGCGCGCGCAGACCAG

Bigger example

```
1 SCGCGCGCGCAGACCAGTTTC
2 ACCAGTTTC$
   AGACCAGTTTC$
   AGTTTCS
   CŚ
6 CAGACCAGTTTC$
   CAGTTTC$
  CCAGTTTC$
   CGCAGACCAGTTTC$
  CGCGCAGACCAGTTTC$
   CGCGCGCAGACCAGTTTC'S G
  CGCGCGCGCAGACCAGTTTCS
  GACCAGTTTC$
14 GCAGACCAGTTTC$
   GCGCAGACCAGTTTC$
16 GCGCGCAGACCAGTTTC$
   GCGCGCGCAGACCAGTTTC$C
   GTTTCS
  TC$
  TTCS
```