# CS 364 COMPUTATIONAL BIOLOGY

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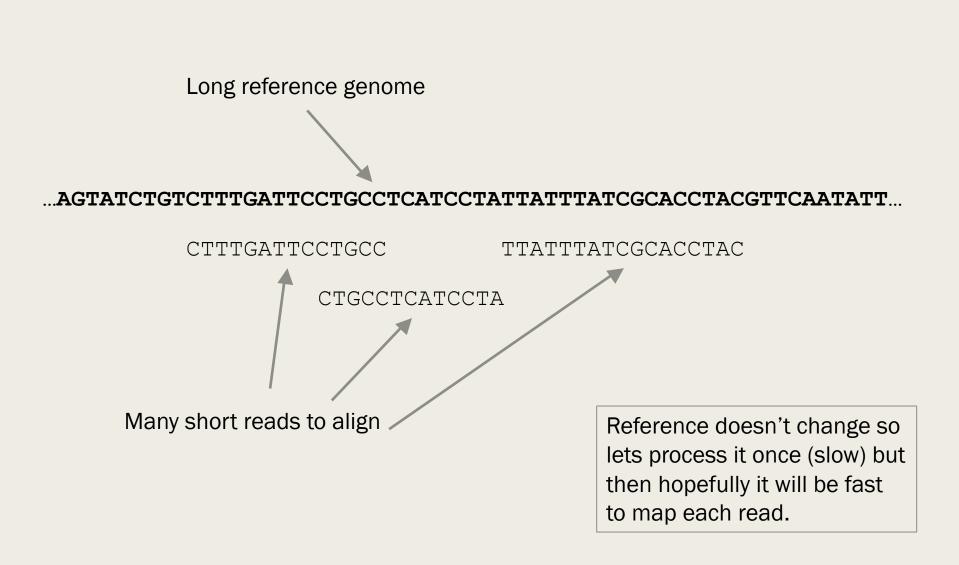
Recap goals of the BWT

FM-Index data structure

Using the BWT for read mapping

# BWT so far...

#### Read mapping



# Bowtie and BWA (posted reading)

■ First practical read aligners to use the Burrows-Wheeler Transform

# Fast and accurate short read alignment with Burrows–Wheeler transform a

Heng Li, Richard Durbin 🖾 💿 Author Notes

*Bioinformatics*, Volume 25, Issue 14, 15 July 2009, Pages 1754–1760, https://doi.org/10.1093/bioinformatics/btp324

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RWA

# Bowtie

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

Ben Langmead 📉, Cole Trapnell, Mihai Pop and Steven L Salzberg

Genome Biology200910:R25https://doi.org/10.1186/gb-2009-10-3-r25©Langmead et al.; licensee BioMed Central Ltd. 2009Received:21 October2008Accepted:4 March 2009Published:4 March 2009Published:4 March 2009

# Comparison of Bowtie and BWA

#### Table 2.

Evaluation on real data

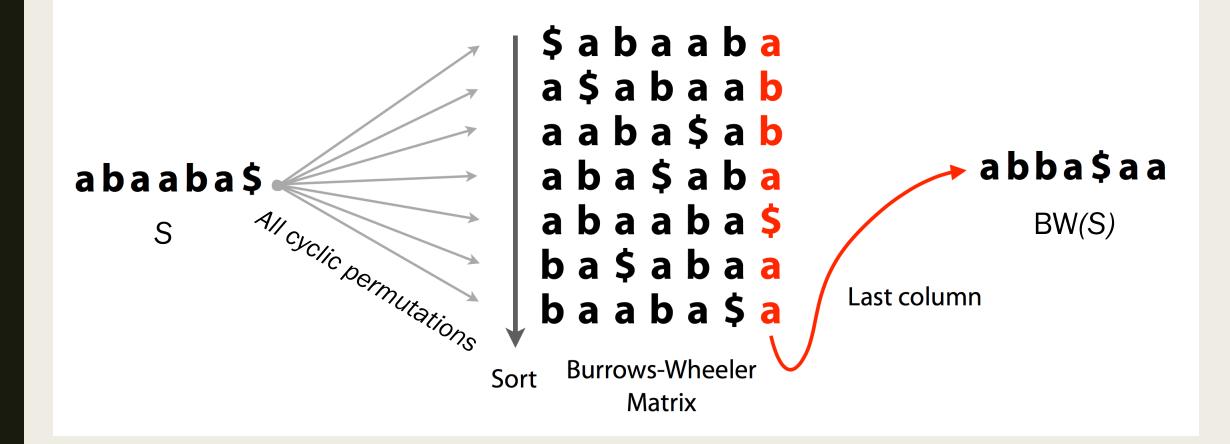
Program	Time (h)	<b>Conf (%)</b>	Paired (%)
Bowtie	5.2	84.4	96.3
BWA	4.0	88.9	98.8
MAQ	94.9	86.1	98.7
SOAP2	3.4	88.3	97.5

e M = # reads naive llionr

The 12.2 million read pairs were mapped to the human genome. CPU time in hours on a single core of a 2.5 GHz Xeon E5420 processor (Time), percent confidently mapped reads (Conf) and percent confident mappings with the mates mapped in the correct orientation and within 300 bp (Paired), are shown in the table.

# BWT

*Reversible permutation* of the characters of a string, used originally for compression



final char (L)	sorted rotations
a	n to decompress. It achieves compression
0	n to perform only comparisons to a depth
0	n transformation} This section describes
0	n transformation} We use the example and
0	n treats the right-hand side as the most
a	n tree for each 16 kbyte input block, enc
a	n tree in the output stream, then encodes
i	n turn, set $L[i]$ to be the
i	n turn, set \$R[i]\$ to the
0	n unusual data. Like the algorithm of Man
a	n use a single set of probabilities table
е	n using the positions of the suffixes in
i	n value at a given point in the vector \$R
е	n we present modifications that improve t
е	n when the block size is quite large. Ho
i	n which codes that have not been seen in
i	n with \$ch\$ appear in the {\em same order
i	n with \$ch\$. In our exam
0	n with Huffman or arithmetic coding. Bri
0	n with figures given by Bell~\cite{bell}.

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

Figure 1: Example of sorted rotations. Twenty consecutive rotations from the sorted list of rotations of a version of this paper are shown, together with the final character of each rotation.

### Compression with BWT

Text: I stuffed a shirt or two into my old carpet-bag, tucked it under my arm, and started for Cape Horn and the Pacific. Quitting the good city of old Manhatto, I duly arrived in New Bedford. It was a Saturday night in December. Much was I disappointed upon learning that the little packet for Nantucket had already sailed, and that no way of reaching that place would offer, till the following Monday. As most young candidates for the pains and penalties of whaling stop at this same New Bedford, thence to embark on their voyage, it may as well be related that I, for one, had no idea of so doing. For my mind was made up to sail in no other than a Nantucket craft, because there was a fine, boisterous something about everything connected with that famous old island, which amazingly pleased me. Besides though New Bedford has of late been gradually monopolising the business of whaling, and though in this matter poor old Nantucket is now much behind her, yet Nantucket was her great original-the Tyre of this Carthage

#### BWT:

e.www.rsn.es,t.dg.ardthene.aenssdgdhs,,n,yyypl,eh,egddgtIoIottaedt,senrt,ds dotdlhotd,dfnesstdyyswrroydntssyyaessdfryskrttoeedtyrfeoysafsdgIrdhdggdglgt rn,efntsd,ep,rtedrhtetdosff,de,tomdgteIeedrgdrtyegdrc

lcMNNNNCsb ec tCwwwwwh ehhhehh e lePphhmrerbhyssphhu nn sfh lldmhScmdwdm-m m a i enaeiuuaa uuauuu ellealoerneleealnnennneneernrirnainieeen a agpmhhthhhhrlhbscdrcnngmdrrllrbDnftvkstttlBBBbbhrw cephhdthfhbhtvtidBnykkkkkpm NNNoooooo dddannnnnnnnanaauuiin gccgtc twwt ttttnttttttt t ffuoi twettcstttttgfhd stcnraat ghmfswhhlhnstlozao ehhh hdl o wulcrrcccccclilps eoooout piaao eioalaaulgar aeea iiareoioieieiaaaaaiaouaioniiiuiiiiiiiia ro  $\cap$ 0 iiiaaaastnwtntnto dpb pfsp M cmqptnffFoff fff Hm hhwymrbnlvou a r pou aooeeoooeeooieeegcooouye lg oraaoaeaaiaaeiaaaneaaiAiaesuiuea iua euii eo i aeeuaeeheiaraaaaiIesfesaranctsair o ev l t.t. n tsaiia nnnnsa idmMttttttooOod o tooaboie eeeo οt mlamlatamldao Tral

### Compression with BWT

- We computed the BWT, and showed that we can reconstruct S from BWT(S)
- BWT(S) is the same length as S, but in practice we can compress it to make it smaller e.g:

S= "Tomorrow\_and\_tomorrow\_and\_tomorrow\$"
BWT(S) = "w\$wwdd\_\_nnoooaattTmmmrrrrrooo\_\_ooo"
Comp = "w\$w2d2 2n2o3a2t2Tm3r6o3 2o3"

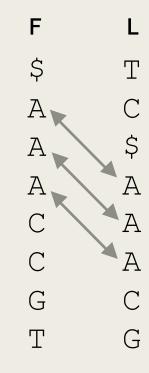
■ In fact, BWT was originally developed as a compression algorithm.

S=ACAACGT

	F		L	
8:	\$.	ACAACG	Т	
3:	A.	ACGT\$A	С	
1:	Α	CAACGT	\$	
4:	Α	CGT\$AC	Α	
2:	C.	AACGT\$	A	
5:	С	GT\$ACA	Α	
6:	G	Г\$АСАА	С	
7:	Τ	\$ACAAC	G	

Τ		\$
С		А
\$	If we have L, i.e.	А
А	BWT(S), we can	Α
А	get F just by	С
А	sorting it $\rightarrow$	С
С		G
G		Т

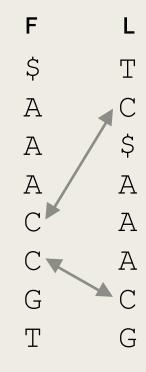
S=ACAACGT\$



Key property 1: If F[i]=X and L[i]=Y, then the string YX must appear in S

Key property 2: The order of each A (also C,G,T) in F is the same as in L

S=ACAACGT\$

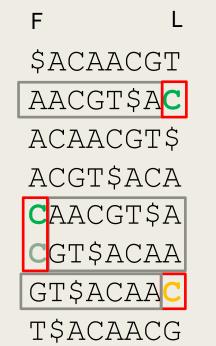


Key property 1: If F[i]=X and L[i]=Y, then the string YX must appear in S

Key property 2: The order of each A (also C,G,T) in F is the same as in L

Key property 2: The order of each C (also each A, G, T) in F is the same as in L

#### S=acaacgt\$



The Cs in the F column are sorted according to the parts of the string that follow them

The Cs in the L column are also sorted according to the parts of the string that follow them

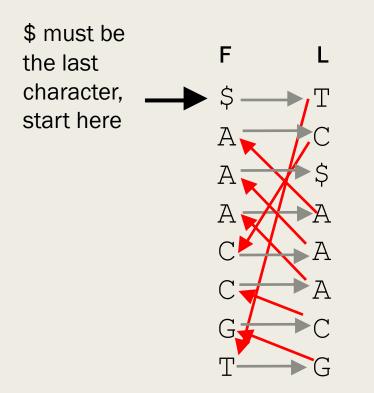
Key property 2: The order of each A (also C, G, T) in F is the same as in L

#### S=ACAACGT\$

#### F

\$ACAACGT AACGT\$AC ACAACGT\$ ACGT\$ACA CAACGT\$A CGT\$ACAA GT\$ACAAC The As in the F column are sorted according to the parts of the string that follow them

The As in the L column are also sorted according to the parts of the string that follow them



Key property 1: If F[i]=X and L[i]=Y, then the string YX must appear in S

Key property 2: The order of each A (also C, G, T) in F is the same as in L

S=...T\$
S=...GT\$
S=...CGT\$
S=...ACGT\$
S=...AACGT\$
S=...CAACGT\$
S=...CAACGT\$
S=...ACAACGT\$

claim, if t has k copies Of char c: c, cz...ck, their order is persevued in L Ti = substring  $\nabla_{1} \leq \nabla_{2} \leq \cdots \leq \nabla_{k} \langle \rangle = \int \nabla_{1} \langle \langle \sigma_{2} \rangle \langle \sigma_{k} \rangle \langle$ A CZ 5 e Tz 215 J2 CK JK 2 2  $\nabla_{k}$ 

# Finish Handout 2

Handout 2 #2 repetitives  $\bigcirc$  $\bigcirc$  $\bigcirc$ 6 X Stop P 7 8 9

# FM-Index data structure

# **FM-Index: data structure for pattern matching**

- Set of auxiliary data structures computed from the BWT of a string S
- The FM-Index consists of 3 parts:

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- The FM-Index consists of 3 parts:

(a) The BWT of S, i.e. the L column of  $\pi^{\text{sorted}}(S)$ 

(b) M[c], the first index of c in F (note that F is actually not part of the FM-Index)

(c) occ(c, i), the number of times c occurs in  $L[1 \cdots i]$ , inclusive

# **FM-Index: data structure for pattern matching**

- Set of auxiliary data structures computed from the BWT of a string S
- The FM-Index consists of 3 parts:

(a) The BWT of S, i.e. the L column of  $\pi^{\text{sorted}}(S)$ 

(b) M[c], the first index of c in F (note that F is actually not part of the FM-Index) (c) occ(c, i), the number of times c occurs in  $L[1 \cdots i]$ , inclusive

- The suffix array A is not technically part of the FM-Index, but we will need it for the last step of finding out where pattern P occurs in the original string S
- A[i] is the index of F[i] in the original string

i	F	L	A	occ(\$)	occ(a)	occ(b)
1	\$ <sub>1</sub> abaab					
2	a <sub>1</sub> \$abaa a <sub>2</sub> aba\$a	b <sub>1</sub>				
3	a <sub>2</sub> aba\$a	b <sub>2</sub>				
4	a <sub>3</sub> ba\$ab a <sub>4</sub> baaba	a <sub>2</sub>				
5	a <sub>4</sub> baaba	\$ <sub>1</sub>				
6	b <sub>1</sub> a\$aba	a <sub>3</sub>				
7	b <sub>2</sub> aaba\$	a <sub>4</sub>				

Step 1: compute the BWT of S

i	F	L	А	occ(\$)	occ(a)	occ(b)
1	\$ <sub>1</sub> abaab	a <sub>1</sub>	7			
2	a <sub>1</sub> \$abaa	$b_1$	6			
3	a <sub>2</sub> aba\$a	$b_2$	3			
4	a <sub>3</sub> ba\$ab	a <sub>2</sub>	4			
5	a <sub>4</sub> baaba	\$ <sub>1</sub>	1			
6	b <sub>1</sub> a\$aba	a <sub>3</sub>	5			
7	b <sub>2</sub> aaba\$	a <sub>4</sub>	2			
			7			

Step 2: compute the suffix array, where A[i] = index of F[i] in the original sequence

i	F		L	Α	occ(\$)		5)	occ(a)	occ(b)
1	\$1	abaab	a <sub>1</sub>	7		0			
2	a <sub>1</sub>	\$abaa	$b_1$	6		0			
3	a <sub>2</sub>	aba\$a	$b_2$	3		0			
4	a <sub>3</sub>	ba\$ab	a <sub>2</sub>	4		0			
5	a <sub>4</sub>	baaba	\$1	1		1			
6	$b_1$	a\$aba	a <sub>3</sub>	5		1			
7	$b_2$	aaba\$	a <sub>4</sub>	2		1			
						4	_		

Step 3: compute the occurrence table for each character c (# times c in L[1...i])

i	F	L	Α	occ(\$)	occ(a	) occ(b)
1	\$ <sub>1</sub> abaab	a <sub>1</sub>	7	0	1	
2	a <sub>1</sub> \$abaa	$b_1$	6	0	1	
3	a <sub>2</sub> aba\$a	$b_2$	3	0	1	
4	a <sub>3</sub> ba\$ab	a <sub>2</sub>	4	0	2	
5	a <sub>4</sub> baaba	\$1	1	1	2	
6	b <sub>1</sub> a\$aba	a <sub>3</sub>	5	1	3	
7	b <sub>2</sub> aaba\$	a <sub>4</sub>	2	1	4	
					~	

Step 3: compute the occurrence table for each character c (# times c in L[1...i])

i	F	L	Α	occ(\$)	occ(a)	0	cc(b	)
1	\$ <sub>1</sub> abaab	a <sub>1</sub>	7	0	1		0	
2	a <sub>1</sub> \$abaa	$b_1$	6	0	1		1	
3	a <sub>2</sub> aba\$a	$b_2$	3	0	1		2	
4	a <sub>3</sub> ba\$ab	a <sub>2</sub>	4	0	2		2	
5	a <sub>4</sub> baaba	\$1	1	1	2		2	
6	b <sub>1</sub> a\$aba	a <sub>3</sub>	5	1	3		2	
7	b <sub>2</sub> aaba\$	a <sub>4</sub>	2	1	4		2	

Step 3: compute the occurrence table for each character c (# times c in L[1...i])

		K	-	_				
	i	F		L	Α	occ(\$)	occ(a)	occ(b)
	1	\$1	abaab a	$a_1$	7	0	1	0
sp(a) —	<b>→</b> 2	a <sub>1</sub>	\$abaa k	01	6	0	1	1
	3	a <sub>2</sub>	aba\$a k	0 <sub>2</sub>	3	0	1	2
	4	a <sub>3</sub>	ba\$ab a	$a_2$	4	0	2	2
ep(a) —	<b>&gt;</b> 5	$a_4$	baaba 4	\$ <sub>1</sub>	1	1	2	2
	6	$b_1$	a\$aba a	a <sub>3</sub>	5	1	3	2
	7	$b_2$	aaba\$ a	$a_4$	2	1	4	2

Step 4: for pattern *P*, start with its last char and compute the start and end points

i	F L	A	occ(\$)	occ(a)	occ(b)
1	$\$_1$ abaab $a_1$	7	0	1	0
2	a <sub>1</sub> \$abaa b <sub>1</sub>	6	0	1	1
3	a <sub>2</sub> aba\$a b <sub>2</sub>	3	0	1	2 🌙
4	$a_3$ ba\$ab $a_2$	4	0	2	2
5	a <sub>4</sub> baaba \$ <sub>1</sub>	1	1	2	2
6	b <sub>1</sub> a\$aba a <sub>3</sub>	5	1	3	2
7	b <sub>2</sub> aaba\$ a <sub>4</sub>	2	1	4	2

0 -> 2 means we must have seen  $b_1$  and  $b_2$  in the L column

i	F L	Α	occ(\$)	occ(a)	occ(b)
1	$a_1$ abaab $a_1$	7	0	1	0
2	a <sub>1</sub> \$abaa b <sub>1</sub>	6	0	1	1
3	$a_2$ abas $a_2b_2$	3	0	1	2 🗸
4	$a_3$ bapap $a_2$	4	0	2	2
5	$a_4$ /a/ba $\$_1$	1	1	2	2
6	b <sub>1</sub> saba a <sub>3</sub>	5	1	3	2
7	b2 aaba\$ a4	2	1	4	2

Find where  $b_1$  and  $b_2$ are in the F column, and repeat the process

	i	F		L	Α	occ(\$)	occ(a)	occ(b)
	1	\$ <sub>1</sub>	abaab	a <sub>1</sub>	7	0	1	0
	2	a <sub>1</sub>	\$abaa	$b_1$	6	0	1	1
	3	a <sub>2</sub>	aba\$a	b	3	0	1	2
	4	a <sub>3</sub>	ba\$ab	a <sub>2</sub>	4	0	2	2
	5	a <sub>4</sub>	Laba	\$1	1	1	2	2
sp(ba)	▶ 6	$b_1$	a\$aba	a <sub>3</sub>	5	1	3	2
ep(ba)-	▶ 7	$b_2$	aaba\$	a <sub>4</sub>	2	1	4	2

i	F	L	Α	occ(\$)	occ(a)	occ(b)
1	\$ <sub>1</sub> abaab	a <sub>1</sub>	7	0	1	0
2	a <sub>1</sub> \$abaa	$b_1$	6	0	1	1
3	a <sub>2</sub> aba\$a	b <sub>2</sub>	3	0	1	2
4	a <sub>3</sub> ba\$ab	a <sub>2</sub>	4	0	2	2
5	a <sub>4</sub> baaba	\$1	1	1	2	2
6	b <sub>1</sub> a\$aba	a <sub>3</sub>	5	1	3	2
7	b <sub>2</sub> aaba\$	a <sub>4</sub>	2	1	4	2

2 -> 4 means we must have seen  $a_3$  and  $a_4$  in the L column

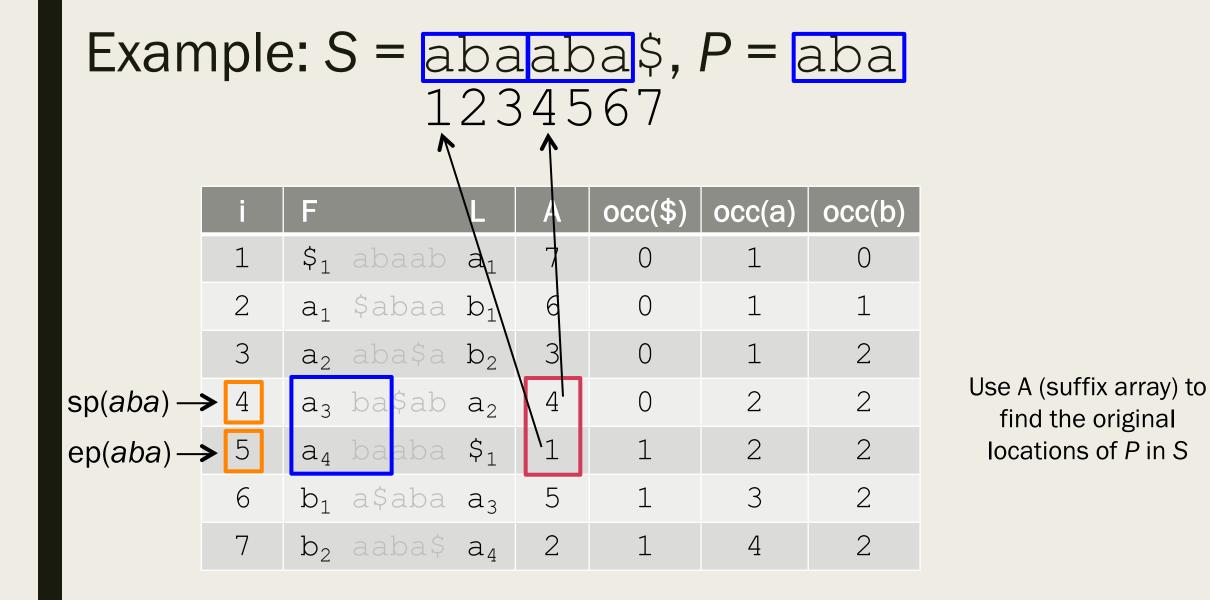
i	F	L	Α	occ(\$)	occ(a)	occ(b)
1	\$ <sub>1</sub> abaab	a <sub>1</sub>	7	0	1	0
2	a <sub>1</sub> \$abaa	b <sub>1</sub>	6	0	1	1
3	a <sub>2</sub> aba\$a	b <sub>2</sub>	3	0	1	2
4	a <sub>3</sub> ba\$ab	a <sub>2</sub>	4	0	2	2
5	a4 baba	\$1	1	1	2	2
6	b <sub>1</sub> as ba	a <sub>3</sub> )	5	1	3	2
7	b <sub>2</sub> aabas	a <sub>4</sub>	2	1	4	2

Find where  $a_3$  and  $a_4$ are in the F column, done since *P* ended

	i	F	L	Α	occ(\$)	occ(a)	occ(b)
	1	\$ <sub>1</sub> abaab	a <sub>1</sub>	7	0	1	0
	2	a <sub>1</sub> \$abaa	$b_1$	6	0	1	1
	3	a <sub>2</sub> aba\$a	$b_2$	3	0	1	2
sp(aba) —	► 4	a₃ ba\$ab	a <sub>2</sub>	4	0	2	2
ep(aba)—	▶ 5	a <sub>4</sub> baaba	\$1	1	1	2	2
	6	b <sub>1</sub> a\$aba	a <sub>3</sub>	5	1	3	2
	7	b <sub>2</sub> aaba\$	a <sub>4</sub>	2	1	4	2

Note that start and end points are inclusive

Step 6: when we reach the end of P, we should have the start/end points in F



Step 7: we are not truly done until we find the locations in the original string!

base case  $sp(\alpha) = 2$ ep(a) = 6 - 1 = 5sp(ba) = 6 + occ(b, sp(a) - 1)occ(b, 2 - 1)= 6 + 0 = 6P(ba) = 6 + Orc(b, PP(a)) - 16+1=7

## BWT pattern matching algorithm

<u>Base case</u>: find the start point (sp) and end point (ep) of the *last* character in P (inclusive, so we subtract 1 from the end point):

 $sp(c) = M[c], \quad ep(c) = M[char alphabetically after c] - 1$ 

<u>Recursion</u>:

$$sp(c\sigma) = M[c] + occ(c, sp(\sigma) - 1)$$
$$ep(c\sigma) = M[c] + occ(c, ep(\sigma)) - 1$$

UBL Pattern Malching Algorithm •base case : let c be last char in P · M table SPECJ = MECJ occ table RP[C] = M[next chair alphabetically]-] staut point in F · A column execursive call i let a be next char in P eb[cd] = W[c] + occ(c' eb(d)) - 1eb[cd] = W[c] + occ(c' eb(d)) - 1end point

# Handout 3

i	F		L	Α	occ(\$)	occ(a)	occ(b)	occ(r)	
1	\$1	barbar	$a_1$						Woi
2	$a_1$	\$barba	r <sub>1</sub>						VVOI
3	a <sub>2</sub>	ra\$bar	$b_1$						1)
4	a <sub>3</sub>	rbara\$	$b_2$						
5	$b_1$	ara\$ba	r <sub>2</sub>						2)
6	$b_2$	arbara	\$1						
7	r <sub>1</sub>	a\$barb	a <sub>2</sub>						
8	r <sub>2</sub>	bara\$b	a <sub>3</sub>						

Work with a partner!

- Fill in a column for A as well
- Try to come up with a formula for sp and ep in terms of M and occ

i	F		L	А	occ(\$)	occ(a)	occ(b)	occ(r)
1	\$1	barbar	a <sub>1</sub>	8	0	1	0	0
2	$a_1$	\$barba	r <sub>1</sub>	7	0	1	0	1
3	a <sub>2</sub>	ra\$bar	$b_1$	5	0	1	1	1
4	a <sub>3</sub>	rbara\$	$b_2$	2	0	1	2	1
5	$b_1$	ara\$ba	r <sub>2</sub>	4	0	1	2	2
6	$b_2$	arbara	\$1	1	1	1	2	2
7	r <sub>1</sub>	a\$barb	a <sub>2</sub>	6	1	2	2	2
8	r <sub>2</sub>	bara\$b	a <sub>3</sub>	3	1	3	2	2

i	F		L	А	occ(\$)	occ(a)	occ(b)	occ(r)
1	\$1	barbar	a <sub>1</sub>	8	0	1	0	0
2	$a_1$	\$barba	r <sub>1</sub>	7	0	1	0	1
3	a <sub>2</sub>	ra\$bar	b <sub>1</sub>	5	0	1	1	1
4	a <sub>3</sub>	rbara\$	$b_2$	2	0	1	2	1
5	$b_1$	ara\$ba	r <sub>2</sub>	4	0	1	2	2
6	$b_2$	arbara	\$1	1	1	1	2	2
7	r <sub>1</sub>	a\$barb	a <sub>2</sub>	6	1	2	2	2
8	r <sub>2</sub>	bara\$b	a <sub>3</sub>	3	1	3	2	2

С	M[c]
\$	1
а	2
b	5
r	7

M[c] is the first index of character c in F (Store instead of F)

i	F		L	А	occ(\$)	occ(a)	occ(b)	occ(r)
1	\$1	barbar	a <sub>1</sub>	8	0	1	0	0
2	$a_1$	\$barba	r <sub>1</sub>	7	0	1	0	1
3	a <sub>2</sub>	ra\$bar	b <sub>1</sub>	5	0	1	1	1
4	a <sub>3</sub>	rbara\$	$b_2$	2	0	1	2	1
5	$b_1$	ara\$ba	r <sub>2</sub>	4	0	1	2	2
6	$b_2$	arbara	\$1	1	1	1	2	2
7	r <sub>1</sub>	a\$barb	a <sub>2</sub>	6	1	2	2	2
8	r <sub>2</sub>	bara\$b	a <sub>3</sub>	3	1	3	2	2

С	M[c]
\$	1
а	2
b	5
r	7

M[c] is the first index of character c in F (Store instead of F)

sp(a) = 2ep(a) = 4

i	F	L	А	occ(\$)	occ(a)	occ(b)	occ(r)
1	\$ <sub>1</sub> barbar	a <sub>1</sub>	8	0	1	0	0
2	a <sub>1</sub> \$barba	r <sub>1</sub>	7	0	1	0	1
3	a <sub>2</sub> ra\$bar	.b <sub>1</sub>	5	0	1	1	1
4	a <sub>3</sub> rbara\$	-b <sub>2</sub>	2	0	1	2	1
5	b <sub>1</sub>	r <sub>2</sub>	4	0	1	2	2
6	b <sub>2</sub> <sup>4</sup> arbara	\$1	1	1	1	2	2
7	r <sub>1</sub> a\$barb	a <sub>2</sub>	6	1	2	2	2
8	r <sub>2</sub> bara\$b	a <sub>3</sub>	3	1	3	2	2

С	M[c]
\$	1
а	2
b	5
r	7

M[c] is the first index of character c in F (Store instead of F)

sp(ba) = M[b] + # b's we saw right before the first a
ep(ba) = M[b] + # b's we saw up until the last a

i	F	L	A	occ(\$)	occ(a)	occ(b)	occ(r)
1	\$1	barbar $a_1$	8	0	1	0	0
2	$a_1$	\$barba r <sub>1</sub>	7	0	1	0	1
3	a <sub>2</sub>	ra\$bar_b <sub>1</sub>	5	0	1	1	1
4	a <sub>3</sub>	rbara\$ b <sub>2</sub>	2	0	1	2	1
5	$b_1$	Laroba r <sub>2</sub>	4	0	1	2	2
6	$b_2$	arbara \$ <sub>1</sub>	1	1	1	2	2
7	r <sub>1</sub>	a\$barb $a_2$	6	1	2	2	2
8	r <sub>2</sub>	bara\$b a <sub>3</sub>	3	1	3	2	2

С	M[c]
\$	1
а	2
b	5
r	7

M[c] is the first index of character c in F (Store instead of F)

sp(ba) = 5 + 0ep(ba) = 5 + 2 - 1 (subtract 1 since we are being *inclusive*)

i	F		L	А	occ(\$)	occ(a)	occ(b)	occ(r)
1	\$1	barbar	a <sub>1</sub>	8	0	1	0	0
2	$a_1$	\$barba	r <sub>1</sub>	7	0	1	0	1
3	a <sub>2</sub>	ra\$bar	$b_1$	5	0	1	1	1
4	a <sub>3</sub>	rbara\$	$b_2$	2	0	1	2	1
5	$b_1$	ara\$ba	r <sub>2</sub>	4	0	1	2	2
6	$b_2$	arbara	\$1	1	1	1	2	2
7	r <sub>1</sub>	a\$barb	a <sub>2</sub>	6	1	2	2	2
8	$r_2$	bara\$b	a <sub>3</sub>	3	1	3	2	2

С	M[c]
\$	1
а	2
b	5
r	7

M[c] is the first index of character c in F (Store instead of F)

sp(ba) = 5ep(ba) = 6

i	F		L	А	occ(\$)	occ(a)	occ(b)	occ(r)
1	\$1	barbar	a <sub>1</sub>	8	0	1	0	0
2	$a_1$	\$barba	r <sub>1</sub>	7	0	1	0	1
3	a <sub>2</sub>	ra\$bar	b <sub>1</sub>	5	0	1	1	1
4	a <sub>3</sub>	rbara\$	$b_2$	2	0	1	2	1
5	b <sub>1</sub>	ara\$ba	r <sub>2</sub>	4	0	1	2	2
6	b <sub>2</sub>	arbara	\$1	1	1	1	2	2
7	r <sub>1</sub>	a\$barb	a <sub>2</sub>	6	1	2	2	2
8	r <sub>2</sub>	bara\$b	a <sub>3</sub>	3	1	3	2	2

С	M[c]
\$	1
а	2
b	5
r	7

sp(ba) = 5ep(ba) = 6

Use A to find locations in original string

UBL P=ba Sp(ba) = M[b] + occ(b, sp(a)-1)5 5 5 5 0 5 0 5 ep(ba) - MEb] + occ(b,ep(a)) - 1= 5+1=6 2. runtime => O(m)

## Pattern matching with BWT

- Setup time O(N)
- Search time O(M)
- Storage space O(N)
  - *O*(1) to store *F* (i.e. *M*)
  - *O*(*N*) to store *L* (i.e. *BWT*(S))
  - O(N) to store A
  - $O(N|\Sigma|)$  to store OCC ("checkpointing" extension allows you to store only part of OCC, without increasing complexity).
- Inexact matching can be implemented in a similar way to inexact matching with little extra cost (as long as few mismatches)

# Summary

Algorithm	Setup time	Lookup time	Storage space
Boyer-Moore	O(M)	O(N)	O(M)
k-mer hash table	O(N)	O(M)	O(N)
BWT/FM-index	O(N)	O(M)	O(N)

But, in practice, for the read mapping problem, BWT approaches have turned out to be the most efficient. Almost all sequence data is processed with a program called *bwa* which uses BWT to map.

## Brief history of BWT and read mapping application

- 1994, BWT introduced (as a compression algorithm)
  - Burrows, M. and Wheeler, D.J. (1994) A block-sorting lossless data compression algorithm. *Technical report 124*, Palo Alto, CA, Digital Equipment Corporation.
- 2000, FM-index for fast searching
  - Ferragina, P. and Manzini, G. (2000) Opportunistic data structures with applications. In *Proceedings of the 41st Symposium on Foundations of Computer Science (FOCS 2000)*, IEEE Computer Society, pp. 390–398.
- 2008, *BWT-SW* for sequence alignment
  - Tam, C. K. Wong, S. M. Yiu (2008) Compressed indexing and local alignment of DNA, Bioinformatics 24
- 2009, *Bowtie* for short read alignment (~19,000 citations to date)
  - Langmead, B. Trapnell, C. Pop, M. Salzberg, S. Ultrafast and memory-efficient alignment of short DNA sequences to the human genome. Genome Biology 10:R25
- 2009, *bwa* (~46,000 citations in 2024)
  - Li, H. and Durbin, R. Fast and accurate short read alignment with Burrows– Wheeler transform Bioinformatics 25: 1754–1760