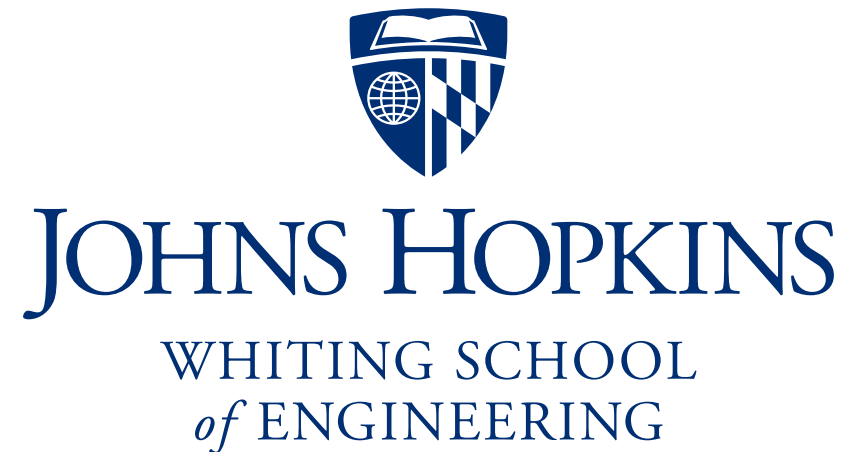


# Burrows-Wheeler Transform and FM Index

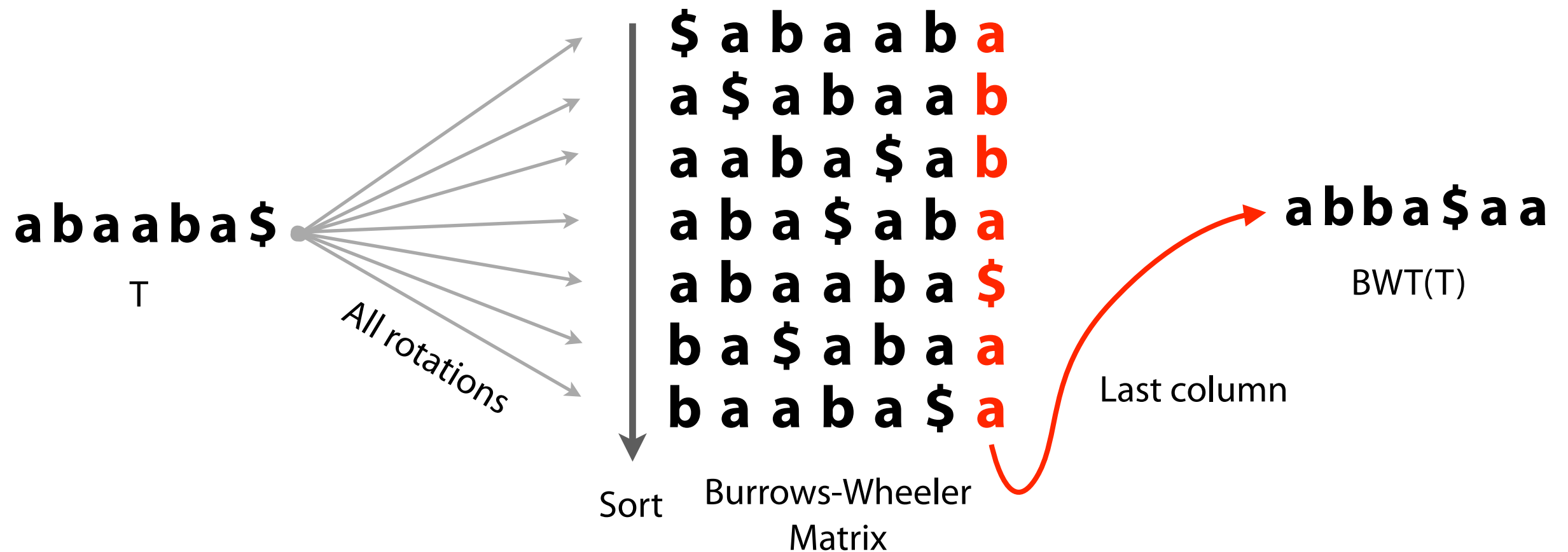
Ben Langmead



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# Burrows-Wheeler Transform

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



How is it useful for compression?

How is it reversible?

How is it an index?

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

# Burrows-Wheeler Transform

```
def rotations(t):  
    """ Return list of rotations of input string t """  
    tt = t * 2  
    return [ tt[i:i+len(t)] for i in xrange(0, len(t)) ]  
  
def bwm(t):  
    """ Return lexicographically sorted list of t's rotations """  
    return sorted(rotations(t))  
  
def bwtViaBwm(t):  
    """ Given T, returns BWT(T) by way of the BWM """  
    return ''.join(map(lambda x: x[-1], bwm(t)))
```

Make list of all rotations

Sort them

Take last column

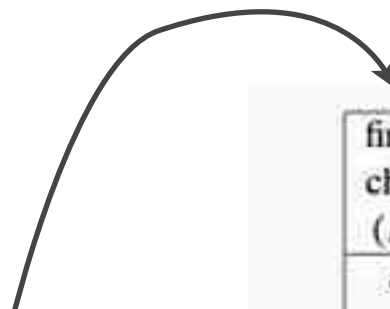
```
>>> bwtViaBwm("Tomorrow_and_tomorrow_and_tomorrow$")  
'w$wwdd__nnoooaattTmmrrrrrrrooo__ooo'  
  
>>> bwtViaBwm("It_was_the_best_of_times_it_was_the_worst_of_times$")  
's$esttssfftteww_hhmmbootttt_ii__woeeaaressi_____  
  
>>> bwtViaBwm('in_the_jingle_jangle_morning_Ill_come_following_you$')  
'u_gleeeengj_mlh1_nnnnt$nwj__lggIolo_iiiiarfcmlylo_oo_'
```

Python example: <http://nbviewer.ipython.org/6798379>

# Burrows-Wheeler Transform

Characters of the BWT are sorted by their *right-context*

This lends additional structure to BWT(T), tending to make it more compressible



final char (L)	sorted rotations
a	n to decompress. It achieves compression
o	n to perform only comparisons to a depth
o	n transformation} This section describes
o	n transformation} We use the example and
o	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
o	n unusual data. Like the algorithm of Man
a	n use a single set of probabilities table
e	n using the positions of the suffixes in
i	n value at a given point in the vector \$R
e	n we present modifications that improve t
e	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
o	n with Huffman or arithmetic coding. Bri
o	n with figures given by Bell~\cite{bell}.

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.

# Burrows-Wheeler Transform

BWM bears a resemblance to the suffix array

\$ a b a a b a  
a \$ a b a a b  
a a b a \$ a b  
a b a \$ a b a  
a b a a b a \$  
b a \$ a b a a  
b a a b a \$ a

BWM(T)

6	\$
5	a \$
2	a a b a \$
3	a b a \$
0	a b a a b a \$
4	b a \$
1	b a a b a \$

SA(T)

Sort order is the same whether rows are rotations or suffixes

# Burrows-Wheeler Transform

In fact, this gives us a new definition / way to construct BWT(T):

$$BWT[i] = \begin{cases} T[SA[i] - 1] & \text{if } SA[i] > 0 \\ \$ & \text{if } SA[i] = 0 \end{cases}$$

“BWT = characters just to the left of the suffixes in the suffix array”

**\$** a b a a b a  
 a **\$** a b a a b  
 a a b a **\$** a b  
 a b a **\$** a b a  
 a b a a b a **\$**  
 b a **\$** a b a a  
 b a a b a **\$** a

BWM(T)

6	<b>\$</b>
5	<b>a</b> <b>\$</b>
2	<b>a</b> <b>a</b> <b>b</b> <b>a</b> <b>\$</b>
3	<b>a</b> <b>b</b> <b>a</b> <b>\$</b>
0	<b>a</b> <b>b</b> <b>a</b> <b>a</b> <b>b</b> <b>a</b> <b>\$</b>
4	<b>b</b> <b>a</b> <b>\$</b>
1	<b>b</b> <b>a</b> <b>a</b> <b>b</b> <b>a</b> <b>\$</b>

SA(T)

# Burrows-Wheeler Transform

```
def suffixArray(s):  
    """ Given T return suffix array SA(T). We use Python's sorted  
        function here for simplicity, but we can do better. """  
    satups = sorted([(s[i:], i) for i in xrange(0, len(s))])  
    # Extract and return just the offsets  
    return map(lambda x: x[1], satups)
```

Make suffix array

```
def bwtViaSa(t):  
    """ Given T, returns BWT(T) by way of the suffix array. """  
    bw = []  
    for si in suffixArray(t):  
        if si == 0: bw.append('$')  
        else: bw.append(t[si-1])  
    return ''.join(bw) # return string-ized version of list bw
```

Take characters just  
to the left of the  
sorted suffixes

```
>>> bwtViaSa("Tomorrow_and_tomorrow_and_tomorrow$")  
'w$wwdd__nnooaaattTmmrrrrrrrooo__ooo'
```

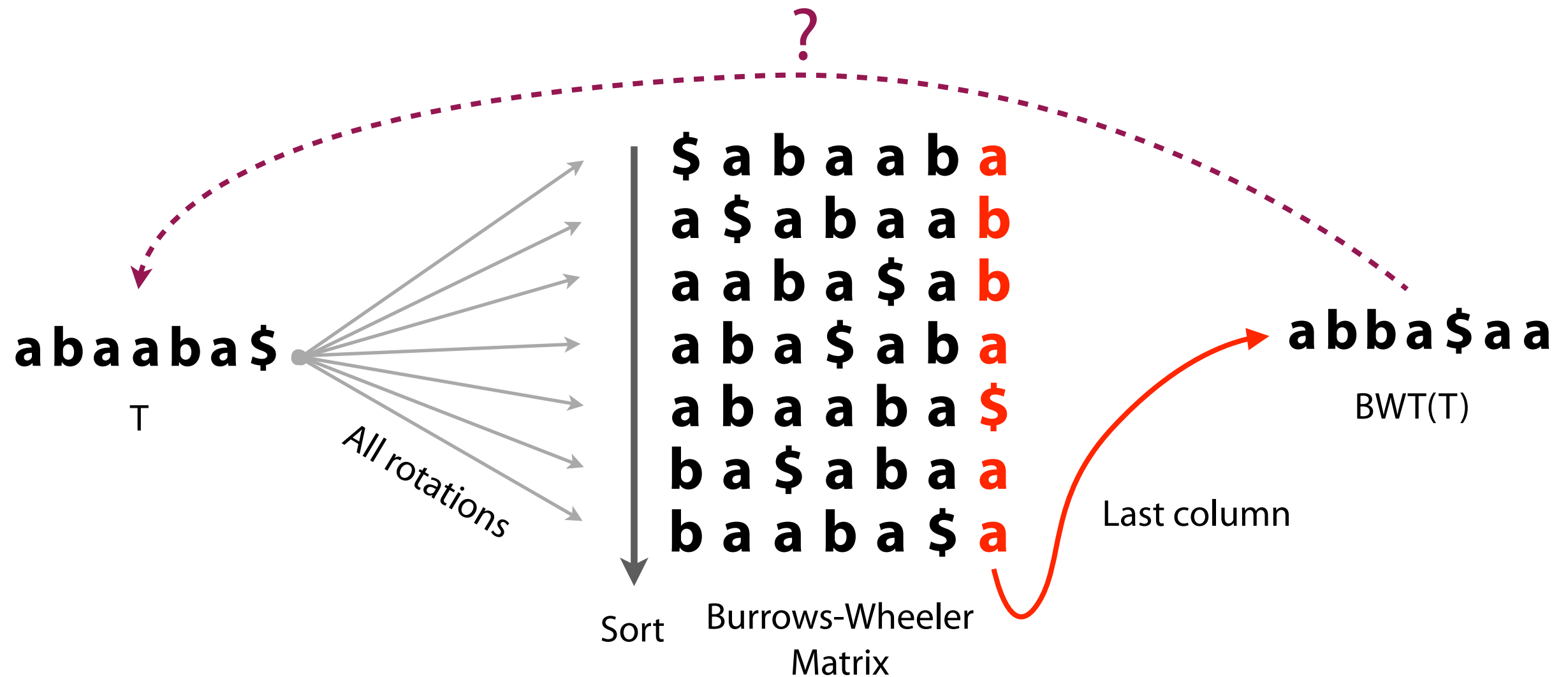
```
>>> bwtViaSa("It_was_the_best_of_times_it_was_the_worst_of_times$")  
's$esttssffteww_hhmmbootttt_ii__woeearessIi_____'
```

```
>>> bwtViaSa('in_the_jingle_jangle_morning_Ill_come_following_you$')  
'u_gleeeengj_mlh1_nnnnt$nwj__lggIolo_iiiiarfcmlylo_oo_'
```

Python example: <http://nbviewer.ipython.org/6798379>

# Burrows-Wheeler Transform

How to reverse the BWT?



BWM has a key property called the *LF Mapping*...



# Burrows-Wheeler Transform: T-ranking

Give each character in  $T$  a rank, equal to # times the character occurred previously in  $T$ . Call this the *T-ranking*.

**a**<sub>0</sub> **b**<sub>0</sub> **a**<sub>1</sub> **a**<sub>2</sub> **b**<sub>1</sub> **a**<sub>3</sub> \$

Now let's re-write the BWM including ranks...

# Burrows-Wheeler Transform

BWM with T-ranking:

	<i>F</i>						<i>L</i>
	\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	<b>a<sub>3</sub></b>
	<b>a<sub>3</sub></b>	\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>
	<b>a<sub>1</sub></b>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>
	<b>a<sub>2</sub></b>	b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>	<b>a<sub>1</sub></b>
	<b>a<sub>0</sub></b>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$
	b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	<b>a<sub>2</sub></b>
	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	<b>a<sub>0</sub></b>

Look at first and last columns, called *F* and *L*

And look at just the **a**s

**a**s occur in the same order in *F* and *L*. As we look down columns, in both cases we see: **a<sub>3</sub>**, **a<sub>1</sub>**, **a<sub>2</sub>**, **a<sub>0</sub>**

# Burrows-Wheeler Transform

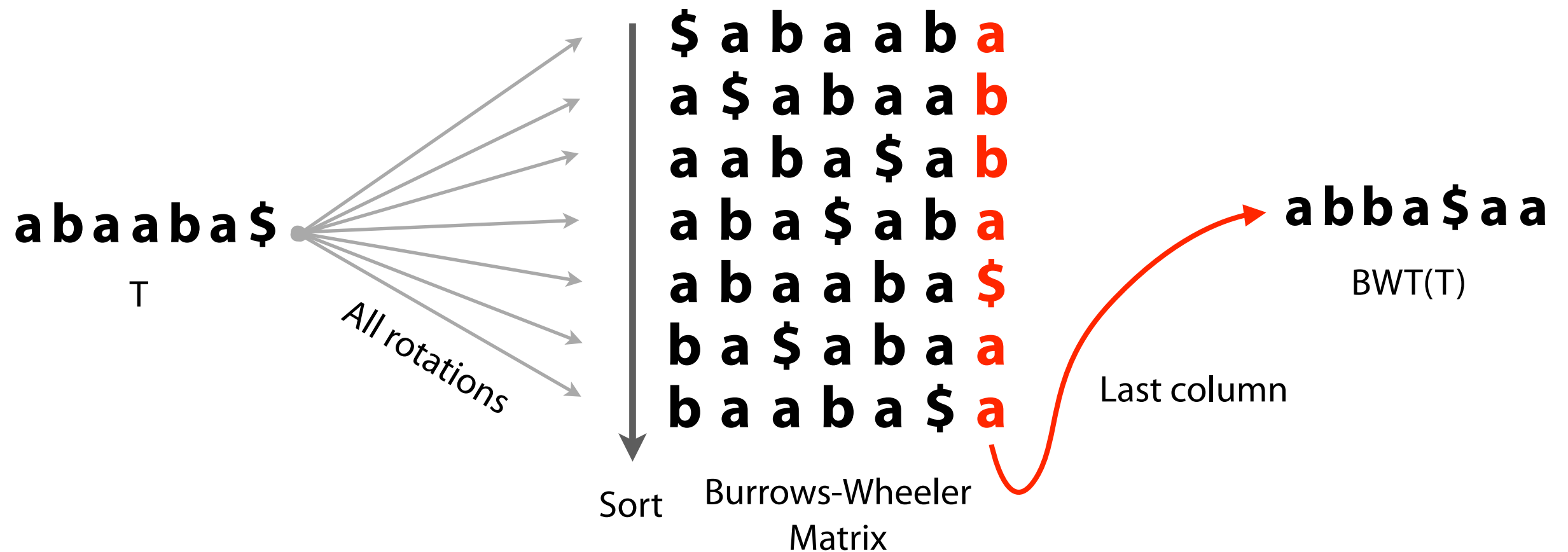
BWM with T-ranking:

<i>F</i>							<i>L</i>
\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	
a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	<b>b<sub>1</sub></b>	
a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	<b>b<sub>0</sub></b>	
a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	
a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	
<b>b<sub>1</sub></b>	a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	
<b>b<sub>0</sub></b>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	

Same with **b**s: **b<sub>1</sub>**, **b<sub>0</sub>**

# Burrows-Wheeler Transform

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



How is it useful for compression?

How is it reversible?

How is it an index?

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

# Burrows-Wheeler Transform: LF Mapping

BWM with T-ranking:

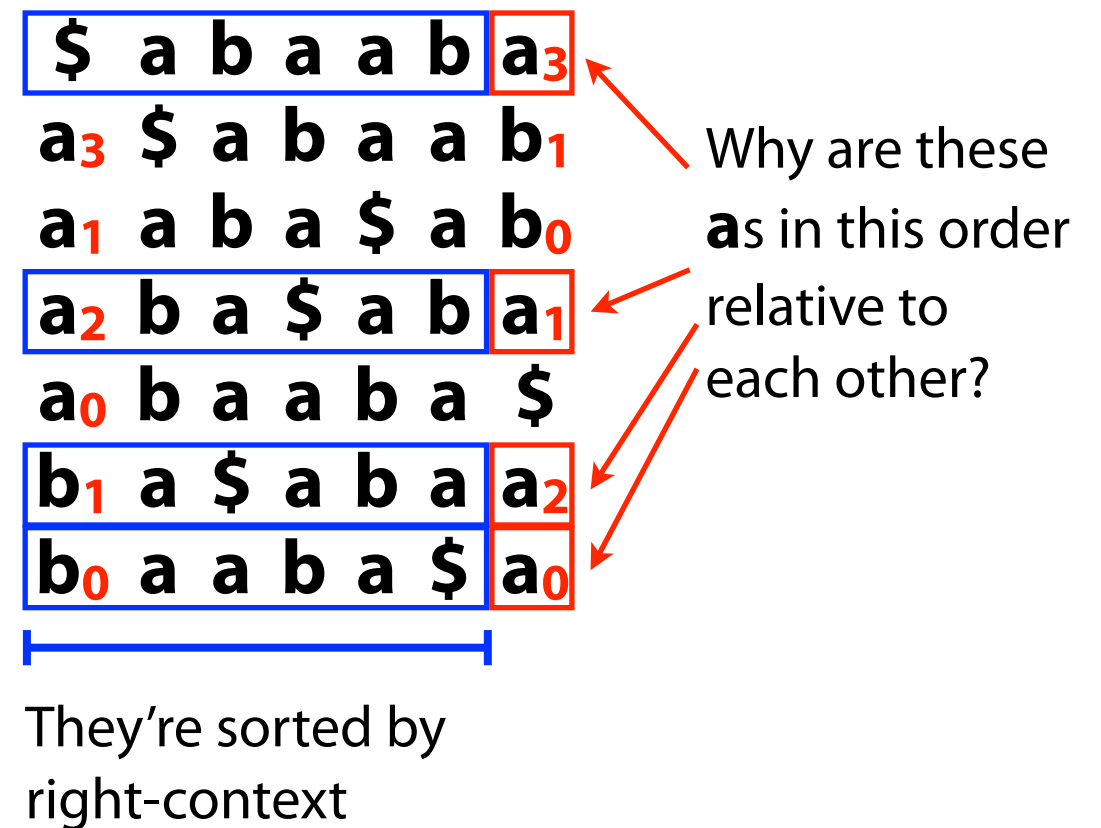
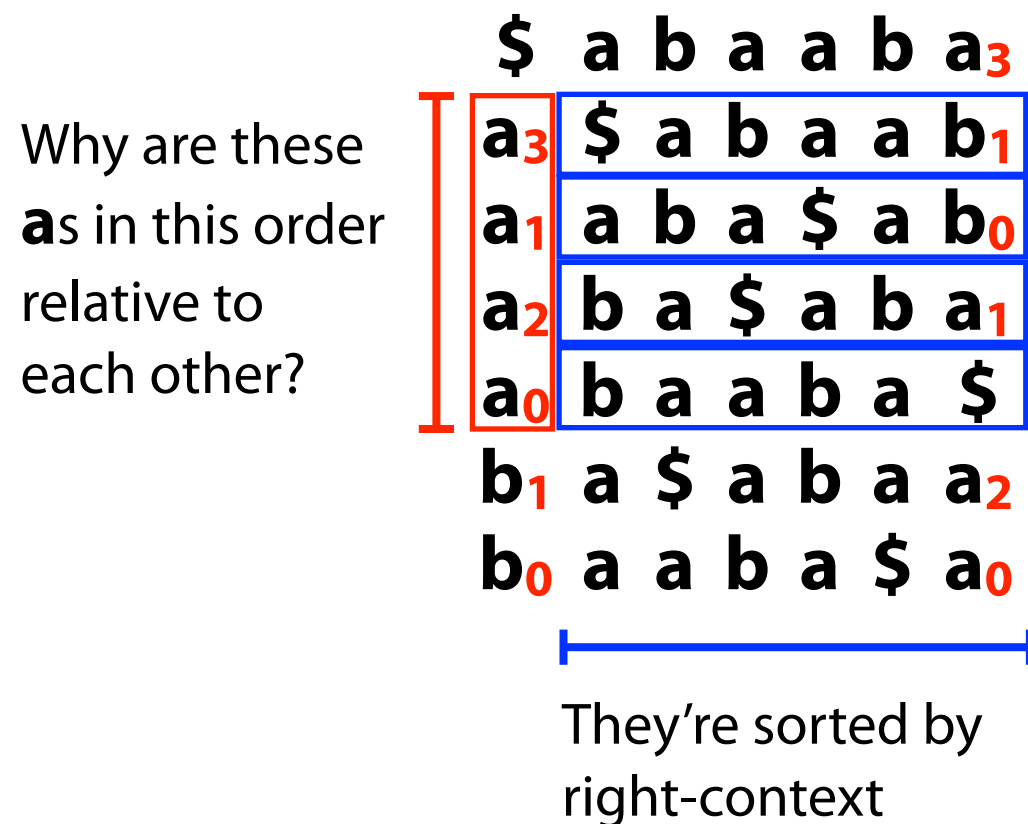
$F$	$L$
\$	<b>a<sub>0</sub></b> <b>b<sub>0</sub></b> <b>a<sub>1</sub></b> <b>a<sub>2</sub></b> <b>b<sub>1</sub></b> <b>a<sub>3</sub></b>
<b>a<sub>3</sub></b>	\$ <b>a<sub>0</sub></b> <b>b<sub>0</sub></b> <b>a<sub>1</sub></b> <b>a<sub>2</sub></b> <b>b<sub>1</sub></b>
<b>a<sub>1</sub></b> <b>a<sub>2</sub></b> <b>b<sub>1</sub></b> <b>a<sub>3</sub></b>	\$ <b>a<sub>0</sub></b> <b>b<sub>0</sub></b>
<b>a<sub>2</sub></b> <b>b<sub>1</sub></b> <b>a<sub>3</sub></b>	\$ <b>a<sub>0</sub></b> <b>b<sub>0</sub></b> <b>a<sub>1</sub></b>
<b>a<sub>0</sub></b> <b>b<sub>0</sub></b> <b>a<sub>1</sub></b> <b>a<sub>2</sub></b> <b>b<sub>1</sub></b> <b>a<sub>3</sub></b>	\$
<b>b<sub>1</sub></b> <b>a<sub>3</sub></b>	\$ <b>a<sub>0</sub></b> <b>b<sub>0</sub></b> <b>a<sub>1</sub></b> <b>a<sub>2</sub></b>
<b>b<sub>0</sub></b> <b>a<sub>1</sub></b> <b>a<sub>2</sub></b> <b>b<sub>1</sub></b> <b>a<sub>3</sub></b>	\$ <b>a<sub>0</sub></b>

LF Mapping: The  $i^{\text{th}}$  occurrence of a character  $c$  in  $L$  and the  $i^{\text{th}}$  occurrence of  $c$  in  $F$  correspond to the *same* occurrence in  $T$

However we rank occurrences of  $c$ , ranks appear in the same order in  $F$  and  $L$

# Burrows-Wheeler Transform: LF Mapping

Why does the LF Mapping hold?



Occurrences of  $c$  in  $F$  are sorted by right-context. Same for  $L$ !

Whatever ranking we give to characters in  $T$ , rank orders in  $F$  and  $L$  will match

# Burrows-Wheeler Transform: LF Mapping

BWM with T-ranking:

<i>F</i>							<i>L</i>
\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	
a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	
a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>	
a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	
a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	
b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	
b <sub>0</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>1</sub>	a <sub>3</sub>	\$	a <sub>0</sub>	

We'd like a different ranking so that for a given character, ranks are in ascending order as we look down the F / L columns...

# Burrows-Wheeler Transform: LF Mapping

BWM with B-ranking:

$F$							$L$
	\$	a <sub>3</sub>	b <sub>1</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>0</sub>	a <sub>0</sub>
	a <sub>0</sub>	\$	a <sub>3</sub>	b <sub>1</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>0</sub>
	a <sub>1</sub>	a <sub>2</sub>	b <sub>0</sub>	a <sub>3</sub>	\$	a <sub>3</sub>	b <sub>1</sub>
	a <sub>2</sub>	b <sub>0</sub>	a <sub>0</sub>	\$	a <sub>3</sub>	b <sub>1</sub>	a <sub>1</sub>
	a <sub>3</sub>	b <sub>1</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>0</sub>	a <sub>0</sub>	\$
	b <sub>0</sub>	a <sub>0</sub>	\$	a <sub>3</sub>	b <sub>1</sub>	a <sub>1</sub>	a <sub>2</sub>
	b <sub>1</sub>	a <sub>1</sub>	a <sub>2</sub>	b <sub>0</sub>	a <sub>0</sub>	\$	a <sub>3</sub>

Ascending rank

$F$  now has very simple structure: a \$, a block of **a**s with ascending ranks, a block of **b**s with ascending ranks



# Burrows-Wheeler Transform

<i>F</i>	<i>L</i>	
\$	<b>a</b> <sub>0</sub>	
<b>a</b> <sub>0</sub>	<b>b</b> <sub>0</sub>	
<b>a</b> <sub>1</sub>	<b>b</b> <sub>1</sub>	← Which BWM row <i>begins</i> with <b>b</b> <sub>1</sub> ?
<b>a</b> <sub>2</sub>	<b>a</b> <sub>1</sub>	Skip row starting with \$ (1 row)
<b>a</b> <sub>3</sub>	\$	Skip rows starting with <b>a</b> (4 rows)
<b>b</b> <sub>0</sub>	<b>a</b> <sub>2</sub>	Skip row starting with <b>b</b> <sub>0</sub> (1 row)
row 6 → <b>b</b> <sub>1</sub>	<b>a</b> <sub>3</sub>	Answer: row 6

# Burrows-Wheeler Transform

Say  $T$  has 300 **A**s, 400 **C**s, 250 **G**s and 700 **T**s and  $\$ < \mathbf{A} < \mathbf{C} < \mathbf{G} < \mathbf{T}$

Which BWM row (0-based) begins with **G**<sub>100</sub>? (Ranks are B-ranks.)

Skip row starting with **\$** (1 row)

Skip rows starting with **A** (300 rows)

Skip rows starting with **C** (400 rows)

Skip first 100 rows starting with **G** (100 rows)

Answer: row  $1 + 300 + 400 + 100 = \mathbf{row\ 801}$

# Burrows-Wheeler Transform: reversing

Reverse BWT( $T$ ) starting at right-hand-side of  $T$  and moving left

**Start** in first row.  $F$  must have  $\$$ .  $L$  contains character just **prior** to  $\$$ : **a<sub>0</sub>**

**a<sub>0</sub>**: LF Mapping says this is same occurrence of **a** as first **a** in  $F$ . **Jump** to row *beginning* with **a<sub>0</sub>**.  $L$  contains character just **prior** to **a<sub>0</sub>**: **b<sub>0</sub>**.

Repeat for **b<sub>0</sub>**, get **a<sub>2</sub>**

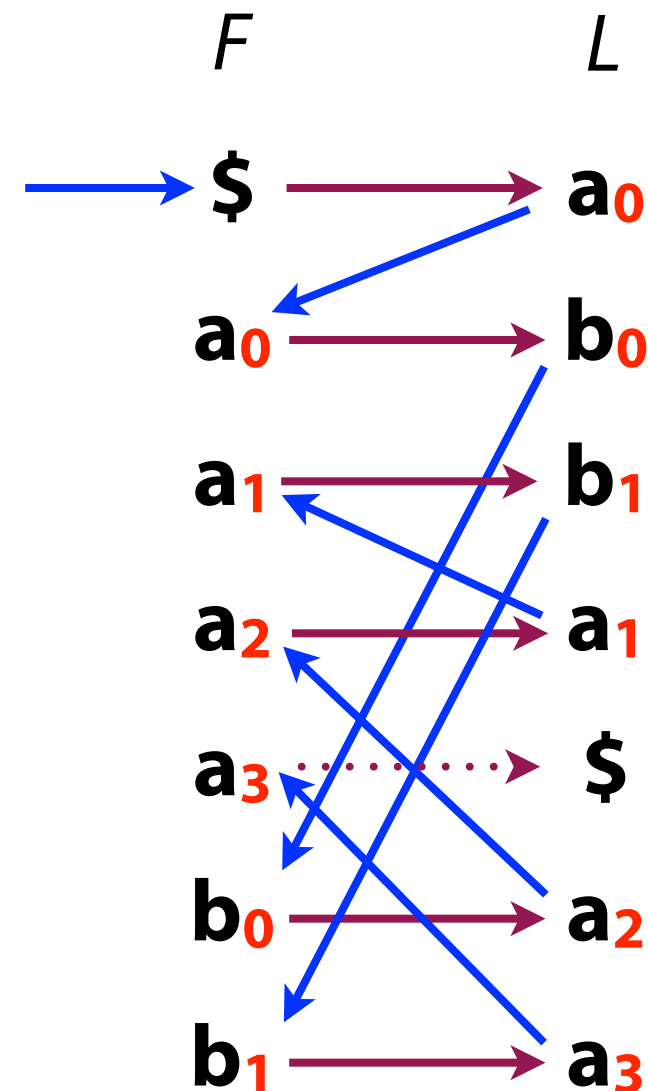
Repeat for **a<sub>2</sub>**, get **a<sub>1</sub>**

Repeat for **a<sub>1</sub>**, get **b<sub>1</sub>**

Repeat for **b<sub>1</sub>**, get **a<sub>3</sub>**

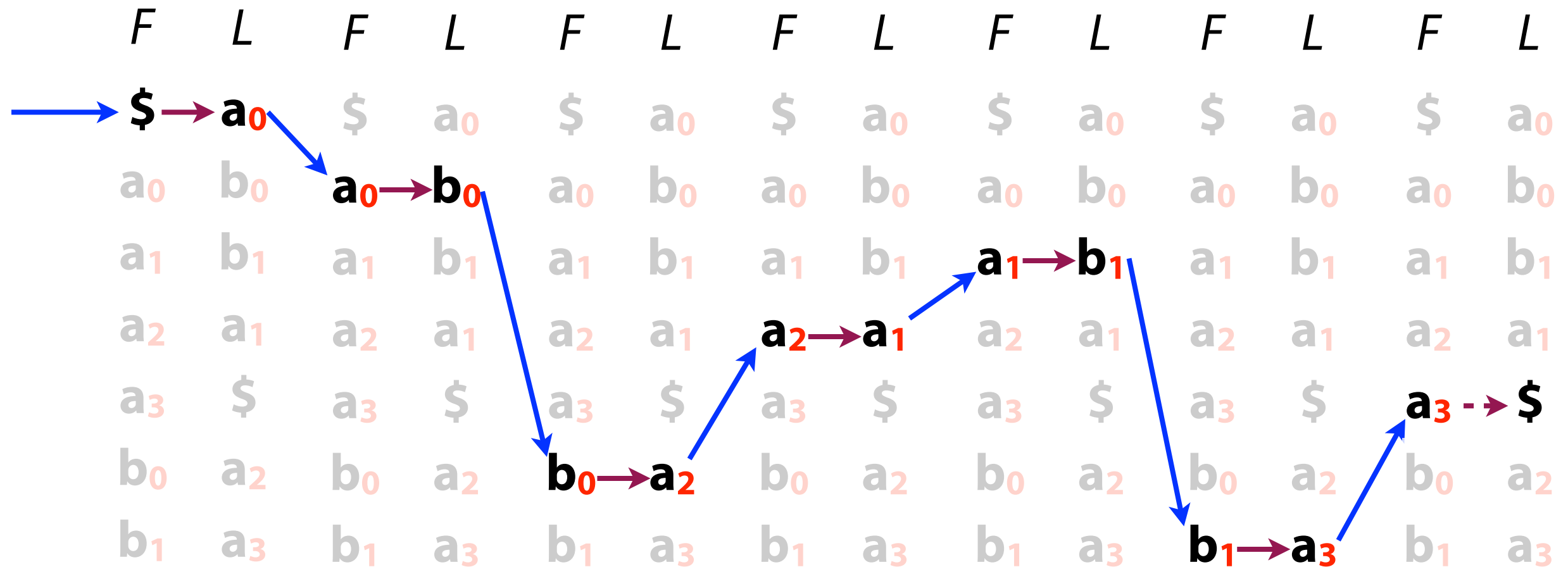
Repeat for **a<sub>3</sub>**, get  $\$$ , done

Reverse of chars we visited = **a<sub>3</sub> b<sub>1</sub> a<sub>1</sub> a<sub>2</sub> b<sub>0</sub> a<sub>0</sub> \$** =  $T$



# Burrows-Wheeler Transform: reversing

Another way to visualize reversing BWT(T):



$T$ :  $a_3 b_1 a_1 a_2 b_0 a_0 \$$

# Burrows-Wheeler Transform: reversing

```
def rankBwt(bw):  
    ''' Given BWT string bw, return parallel list of B-ranks. Also  
        returns tots: map from character to # times it appears. '''  
    tots = dict()  
    ranks = []  
    for c in bw:  
        if c not in tots: tots[c] = 0  
        ranks.append(tots[c])  
        tots[c] += 1  
    return ranks, tots
```

Calculate B-ranks and count  
occurrences of each char

```
def firstCol(tots):  
    ''' Return map from character to the range of rows prefixed by  
        the character. '''  
    first = {}  
    totc = 0  
    for c, count in sorted(tots.iteritems()):  
        first[c] = (totc, totc + count)  
        totc += count  
    return first
```

Make concise representation  
of first BWM column

```
def reverseBwt(bw):  
    ''' Make T from BWT(T) '''  
    ranks, tots = rankBwt(bw)  
    first = firstCol(tots)  
    rowi = 0 # start in first row  
    t = '$' # start with rightmost character  
    while bw[rowi] != '$':  
        c = bw[rowi]  
        t = c + t # prepend to answer  
        # jump to row that starts with c of same rank  
        rowi = first[c][0] + ranks[rowi]  
    return t
```

Do reversal

Python example:

<http://nbviewer.ipython.org/6860491>

# Burrows-Wheeler Transform: reversing

```
>>> reverseBwt("w$wdd__nnooaattTmmrrrrrrrooo__ooo")
'Tomorrow_and_tomorrow_and_tomorrow$'

>>> reverseBwt("s$esttssfftteww_hhmmbootttt_i__woeearessIi_____")
'It_was_the_best_of_times_it_was_the_worst_of_times$'

>>> reverseBwt("u_gleeeengj_mlh1_nnnnt$nwj__lggIolo_iiiarfcmylo_oo_")
'in_the_jingle_jangle_morning_Ill_come_following_you$'
```

ranks list is  $m$  integers  
long! We'll fix later.

```
def reverseBwt(bw):
    ''' Make T from BWT(T) '''
    ranks, tots = rankBwt(bw)
    first = firstCol(tots)
    rowi = 0 # start in first row
    t = '$' # start with rightmost character
    while bw[rowi] != '$':
        c = bw[rowi]
        t = c + t # prepend to answer
        # jump to row that starts with c of same rank
        rowi = first[c][0] + ranks[rowi]
    return t
```

# Burrows-Wheeler Transform

We've seen how BWT is useful for compression:

Sorts characters by right-context, making a more compressible string

And how it's reversible:

Repeated applications of LF Mapping, recreating  $T$  from right to left

How is it used as an index?

# FM Index

FM Index: an index combining the BWT *with a few small auxilliary data structures*

“FM” supposedly stands for “Full-text Minute-space.”  
(But inventors are named Ferragina and Manzini)

Core of index consists of  $F$  and  $L$  from BWM:

$F$  can be represented very simply  
(1 integer per alphabet character)

And  $L$  is compressible

Potentially very space-economical!

$F$							$L$
<b>\$</b>	a	b	a	a	b	a	
<b>a</b>	\$	a	b	a	a	b	
<b>a</b>	a	b	a	\$	a	b	
<b>a</b>	b	a	\$	a	b	a	
<b>a</b>	b	a	a	b	a	\$	
<b>b</b>	a	\$	a	b	a	a	
<b>b</b>	a	a	b	a	\$	a	

└──────────┘  
Not stored in index


Paolo Ferragina, and Giovanni Manzini. "Opportunistic data structures with applications." *Foundations of Computer Science, 2000. Proceedings. 41st Annual Symposium on*. IEEE, 2000.



# FM Index: querying

Though BWM is related to suffix array, we can't query it the same way

\$	a	b	a	a	b	a
a	\$	a	b	a	a	b
a	a	b	a	\$	a	b
a	b	a	\$	a	b	a
a	b	a	a	b	a	\$
b	a	\$	a	b	a	a
b	a	a	b	a	\$	a



6	\$
5	a \$
2	a a b a \$
3	a b a \$
0	a b a a b a \$
4	b a \$
1	b a a b a \$

We don't have these columns; binary search isn't possible



# FM Index: querying

Look for range of rows of BWM(T) with  $P$  as prefix

Do this for  $P$ 's shortest suffix, then extend to successively longer suffixes until range becomes empty or we've exhausted  $P$

$P = \mathbf{ab}\mathbf{a}$

Easy to find all the rows beginning with  $\mathbf{a}$ , thanks to  $F$ 's simple structure

$F$						$L$
\$	a	b	a	a	b	$\mathbf{a}_3$
$\mathbf{a}_0$	\$	a	b	a	a	$\mathbf{b}_1$
$\mathbf{a}_1$	a	b	a	\$	a	$\mathbf{b}_0$
$\mathbf{a}_2$	b	a	\$	a	b	$\mathbf{a}_1$
$\mathbf{a}_3$	b	a	a	b	a	\$
$\mathbf{b}_0$	a	\$	a	b	a	$\mathbf{a}_2$
$\mathbf{b}_1$	a	a	b	a	\$	$\mathbf{a}_0$

# FM Index: querying

We have rows beginning with **a**, now we seek rows beginning with **ba**

$P = \mathbf{aba}$

$P = \mathbf{a} \mathbf{b} \mathbf{a}$

	$F$		$L$
\$	a	b	a
a <sub>0</sub>	\$	a	b
a <sub>1</sub>	a	b	a
a <sub>2</sub>	b	a	\$
a <sub>3</sub>	b	a	a
b <sub>0</sub>	a	\$	a
b <sub>1</sub>	a	a	b

← Look at those rows in  $L$ .  
 $\mathbf{b}_0, \mathbf{b}_1$  are  $\mathbf{b}$ s occuring just to left.

Use LF Mapping. Let new range delimit those **bs**

	$F$		$L$
	\$	a	b
$a_0$	\$	a	b
$a_1$	a	b	\$
$a_2$	b	a	\$
$a_3$	b	a	a
$b_0$	a	\$	a
$b_1$	a	a	b

Now we have the rows with prefix **ba**

# FM Index: querying

We have rows beginning with **ba**, now we seek rows beginning with **aba**

$P = \mathbf{aba}$

$F$							$L$
\$	a	b	a	a	b		$a_0$
$a_0$	\$	a	b	a	a		$b_0$
$a_1$	a	b	a	\$	a		$b_1$
$a_2$	b	a	\$	a	b		$a_1$
$a_3$	b	a	a	b	a		\$
$b_0$	a	\$	a	b	a		$a_2$
$b_1$	a	a	b	a	\$		$a_3$

←  $a_2, a_3$  occur just to left.

Use LF Mapping →

$P = \mathbf{aba}$

$F$							$L$
\$	a	b	a	a	b		$a_0$
$a_0$	\$	a	b	a	a		$b_0$
$a_1$	a	b	a	\$	a		$b_1$
$a_2$	b	a	\$	a	b		$a_1$
$a_3$	b	a	a	b	a		\$
$b_0$	a	\$	a	b	a		$a_2$
$b_1$	a	a	b	a	\$		$a_3$

Now we have the rows with prefix **aba**

# FM Index: querying

$P = \text{aba}$

Now we have the same range,  $[3, 5)$ , we would have got from querying suffix array

$F$		$L$
\$	a b a a b	<b>a<sub>0</sub></b>
<b>a<sub>0</sub></b>	\$ a b a a	<b>b<sub>0</sub></b>
<b>a<sub>1</sub></b>	a b a \$ a	<b>b<sub>1</sub></b>
$[3, 5)$	<b>a<sub>2</sub></b> b a \$ a b	<b>a<sub>1</sub></b>
	<b>a<sub>3</sub></b> b a a b a	\$
	<b>b<sub>0</sub></b> a \$ a b a	<b>a<sub>2</sub></b>
	<b>b<sub>1</sub></b> a a b a \$	<b>a<sub>3</sub></b>

Where are these?

6	\$
5	a \$
2	a a b a \$
$[3, 5)$	<b>a b a \$</b>
	<b>a b a a b a \$</b>
4	b a \$
1	b a a b a \$

Unlike suffix array, we don't immediately know *where* the matches are in T...

# FM Index: querying

When  $P$  does not occur in  $T$ , we will eventually fail to find the next character in  $L$ :

$$P = \mathbf{bba}$$

$F$							$L$
\$	a	b	a	a	b	a	$a_0$
$a_0$	\$	a	b	a	a	b	$b_0$
$a_1$	a	b	a	\$	a	b	$b_1$
$a_2$	b	a	\$	a	b	a	$a_1$
$a_3$	b	a	a	b	a	\$	
Rows with <b>ba</b> prefix							
$b_0$	a	\$	a	b	a	$a_2$	← No <b>bs</b> !
$b_1$	a	a	b	a	\$	$a_3$	

# FM Index: querying

If we *scan* characters in the last column, that can be very slow,  $O(m)$

$P = \mathbf{ab}\mathbf{a}$

$F$						$L$
\$	a	b	a	a	b	<b>a<sub>3</sub></b>
<b>a<sub>0</sub></b>	\$	a	b	a	a	<b>b<sub>1</sub></b>
<b>a<sub>1</sub></b>	a	b	a	\$	a	<b>b<sub>0</sub></b>
<b>a<sub>2</sub></b>	b	a	\$	a	b	<b>a<sub>1</sub></b>
<b>a<sub>3</sub></b>	b	a	a	b	a	\$
<b>b<sub>0</sub></b>	a	\$	a	b	a	<b>a<sub>2</sub></b>
<b>b<sub>1</sub></b>	a	a	b	a	\$	<b>a<sub>0</sub></b>

Scan, looking for **b**s

# FM Index: lingering issues

(1) Scanning for preceding character is slow

	\$	a	b	a	a	b	<b>a<sub>0</sub></b>
<b>a<sub>0</sub></b>	\$	a	b	a	a	b	<b>b<sub>0</sub></b>
<b>a<sub>1</sub></b>	a	b	a	\$	a	b	<b>b<sub>1</sub></b>
<b>a<sub>2</sub></b>	b	a	\$	a	b	a	<b>a<sub>1</sub></b>
<b>a<sub>3</sub></b>	b	a	a	b	a	\$	
<b>b<sub>0</sub></b>	a	\$	a	b	a	a	<b>a<sub>2</sub></b>
<b>b<sub>1</sub></b>	a	a	b	a	\$	a	<b>a<sub>3</sub></b>

$O(m)$   
scan

(2) Storing ranks takes too much space

```
def reverseBwt(bw):
    """ Make T from BWT(T) """
    ranks, tots = rankBwt(bw)
    first = firstCol(tots)
    rowi = 0
    t = "$"
    while bw[rowi] != '$':
        c = bw[rowi]
        t = c + t
        rowi = first[c][0] + ranks[rowi]
    return t
```

$m$  integers

(3) Need way to find where matches occur in  $T$ :

Where?

	\$	a	b	a	a	b	<b>a<sub>0</sub></b>
<b>a<sub>0</sub></b>	\$	a	b	a	a	b	<b>b<sub>0</sub></b>
<b>a<sub>1</sub></b>	a	b	a	\$	a	b	<b>b<sub>1</sub></b>
<b>a<sub>2</sub></b>	b	a	\$	a	b	a	<b>a<sub>1</sub></b>
<b>a<sub>3</sub></b>	b	a	a	b	a	\$	
<b>b<sub>0</sub></b>	a	\$	a	b	a	a	<b>a<sub>2</sub></b>
<b>b<sub>1</sub></b>	a	a	b	a	\$	a	<b>a<sub>3</sub></b>



# FM Index: fast rank calculations

Is there an  $O(1)$  way to determine which **b**s precede the **a**s in our range?

<i>F</i>		<i>L</i>
\$	a b a a b	<b>a<sub>0</sub></b>
<b>a<sub>0</sub></b>	\$ a b a a	<b>b<sub>0</sub></b>
<b>a<sub>1</sub></b>	a b a \$ a	<b>b<sub>1</sub></b>
<b>a<sub>2</sub></b>	b a \$ a b	<b>a<sub>1</sub></b>
<b>a<sub>3</sub></b>	b a a b a	\$
<b>b<sub>0</sub></b>	a \$ a b a	<b>a<sub>2</sub></b>
<b>b<sub>1</sub></b>	a a b a \$	<b>a<sub>3</sub></b>

Idea: pre-calculate # **a**s, **b**s in *L* up to every row:

<i>F</i>	<i>L</i>	<i>Tally</i>	
		<b>a</b>	<b>b</b>
\$	<b>a</b>	1	0
<b>a</b>	<b>b</b>	1	1
<b>a</b>	<b>b</b>	1	2
<b>a</b>	<b>a</b>	2	2
<b>a</b>	\$	2	2
<b>b</b>	<b>a</b>	3	2
<b>b</b>	<b>a</b>	4	2

We infer **b<sub>0</sub>** and **b<sub>1</sub>** appear in *L* in this range

$O(1)$  time, but requires  $m \times |\Sigma|$  integers

# FM Index: fast rank calculations

Another idea: pre-calculate # **a**s, **b**s in  $L$  up to *some* rows, e.g. every 5<sup>th</sup> row.  
Call pre-calculated rows *checkpoints*.

		<i>Tally</i>		
$F$	$L$	<b>a</b>	<b>b</b>	
\$	<b>a</b>	1	0	← Lookup here succeeds as usual
<b>a</b>	<b>b</b>			
<b>a</b>	<b>b</b>			
<b>a</b>	<b>a</b>			
<b>a</b>	\$			← Oops: not a checkpoint
<b>b</b>	<b>a</b>	3	2	← But there's one nearby
<b>b</b>	<b>a</b>			

To resolve a lookup for character  $c$  in non-checkpoint row, scan along  $L$  until we get to nearest checkpoint. Use tally at the checkpoint, *adjusted for # of cs we saw along the way*.

# FM Index: fast rank calculations

What's my rank?

$$482 + 2 - 1 = 483$$

Checkpoint      ↑      ↑      ↑  
as along the way      tally -> rank

What's my rank?

$$439 - 2 - 1 = 436$$

*Tally*

<i>L</i>	<b>a</b>	<b>b</b>
⋮	⋮	⋮
a	482	432
b		
b		
a		
<b>a</b>		
a		
a		
b		
b		
<b>b</b>		
a		
a		
b		
b	488	439
a		
b		

Assuming checkpoints are spaced  $O(1)$   
distance apart, lookups are  $O(1)$

# FM Index: a few problems

Solved! At the expense of adding checkpoints ( $O(m)$  integers) to index.

(1)

	$F$		$L$				
	\$	a	b	a	a	b	$a_0$
$a_0$	\$	a	b	a	a	$b_0$	
$a_1$	a	b	a	\$	a	$b_1$	
$a_2$	b	a	\$	a	b	$a_1$	
$a_3$	b	a	a	b	a	\$	
$b_0$	a	\$	a	b	a	$a_2$	
$b_1$	a	a	b	a	\$	$a_3$	

This scan is  $O(m)$  work

With checkpoints it's  $O(1)$

(2) Ranking takes too much space

$m$  integers

```
def reverseBwt(bw):  
    """ Make T from BWT(T) """  
    ranks, tots = rankBwt(bw)  
    first = firstCol(tots)  
    rowi = 0  
    t = "$"  
    while bw[rowi] != '$':  
        c = bw[rowi]  
        t = c + t  
        rowi = first[c][0] + ranks[rowi]  
    return t
```

With checkpoints, we greatly reduce  
# integers needed for ranks

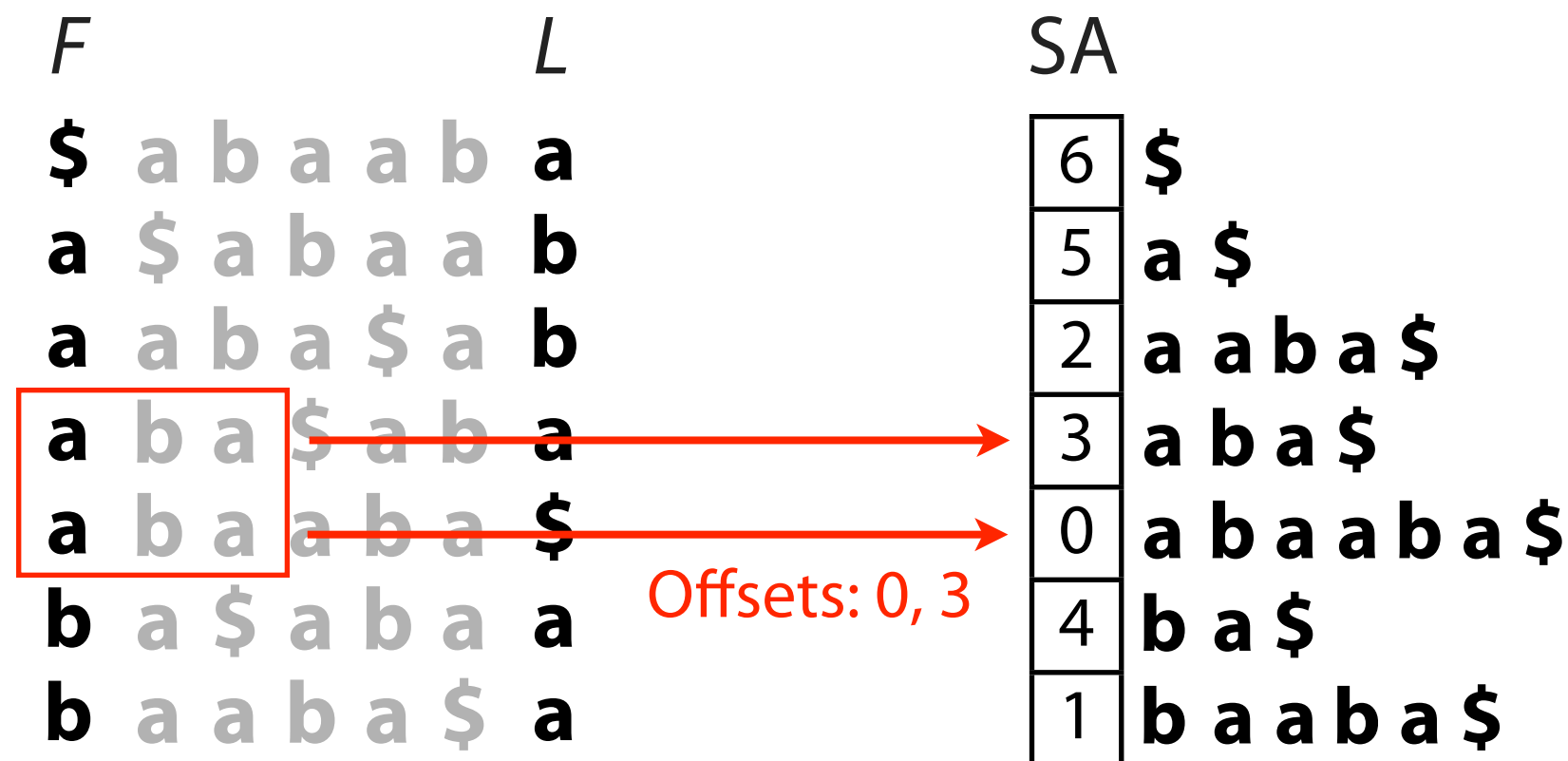
But it's still  $O(m)$  space - there's literature  
on how to improve this space bound

# FM Index: a few problems

Not yet solved: **(3)** Need a way to find where these occurrences are in  $T$ :

\$	a	b	a	a	b	a <sub>0</sub>
a <sub>0</sub>	\$	a	b	a	a	b <sub>0</sub>
a <sub>1</sub>	a	b	a	\$	a	b <sub>1</sub>
a <sub>2</sub>	b	a	\$	a	b	a <sub>1</sub>
a <sub>3</sub>	b	a	a	b	a	\$
b <sub>0</sub>	a	\$	a	b	a	a <sub>2</sub>
b <sub>1</sub>	a	a	b	a	\$	a <sub>3</sub>

If suffix array were part of index, we could simply look up the offsets

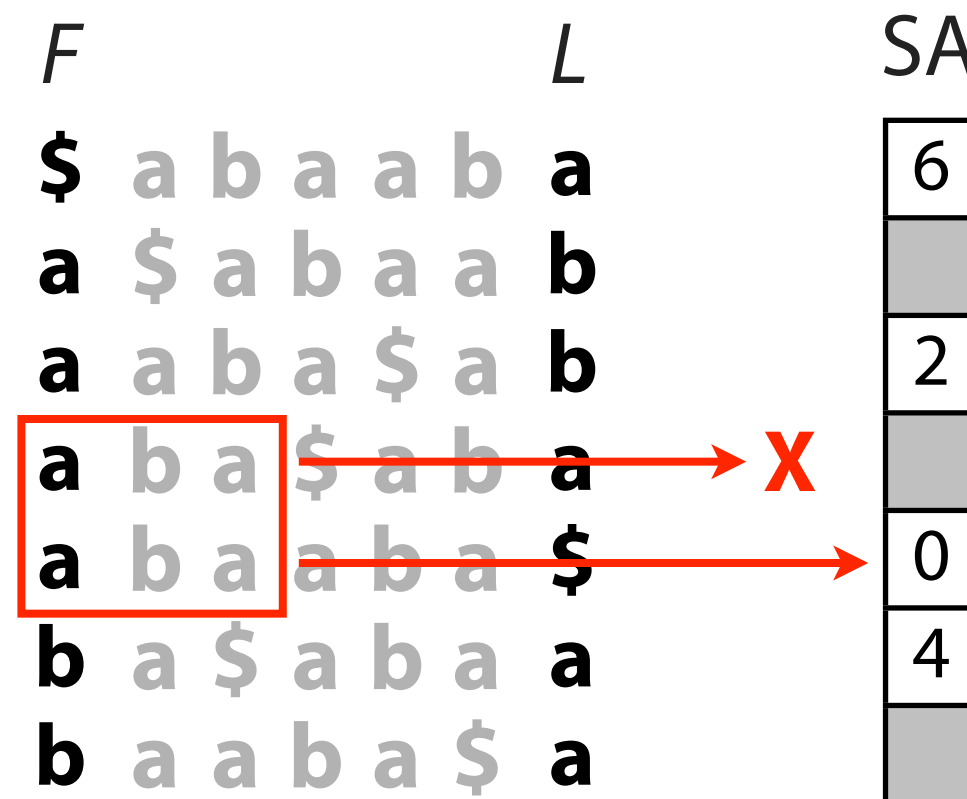


But SA requires  $m$  integers

# FM Index: resolving offsets

Idea: store some, but not all, entries of the suffix array

<i>F</i>						<i>L</i>		<i>SA</i>
\$	a	b	a	a	b	a		6
a	\$	a	b	a	a	b		
a	a	b	a	\$	a	b		2
a	b	a	\$	a	b	a		
a	b	a	a	b	a	\$		0
b	a	\$	a	b	a	a		4
b	a	a	b	a	\$	a		



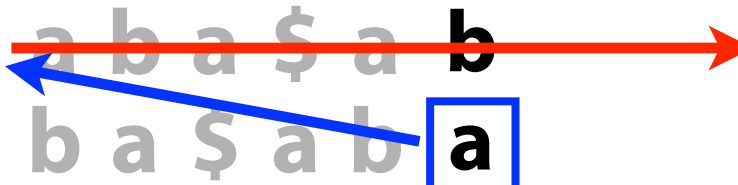
Lookup for row 4 succeeds - we kept that entry of SA

Lookup for row 3 fails - we discarded that entry of SA

# FM Index: resolving offsets

But LF Mapping tells us that the **a** at the end of row 3 corresponds to...  
...the **a** at the beginning of row 2

<i>F</i>							<i>L</i>		<i>SA</i>
<b>\$</b>	<b>a</b>	<b>b</b>	<b>a</b>	<b>a</b>	<b>b</b>	<b>a</b>			6
<b>a</b>	<b>\$</b>	<b>a</b>	<b>b</b>	<b>a</b>	<b>a</b>	<b>b</b>			
<b>a</b>	<b>a</b>	<b>b</b>	<b>a</b>	<b>\$</b>	<b>a</b>	<b>b</b>			2
<b>a</b>	<b>b</b>	<b>a</b>	<b>\$</b>	<b>a</b>	<b>b</b>	<b>a</b>			
<b>a</b>	<b>b</b>	<b>a</b>	<b>a</b>	<b>b</b>	<b>a</b>	<b>\$</b>			0
<b>b</b>	<b>a</b>	<b>\$</b>	<b>a</b>	<b>b</b>	<b>a</b>	<b>a</b>			4
<b>b</b>	<b>a</b>	<b>a</b>	<b>b</b>	<b>a</b>	<b>\$</b>	<b>a</b>			



And row 2 has a suffix array value = 2

So row 3 has suffix array value = 3 = 2 (row 2's SA val) + 1 (# steps to row 2)

If saved SA values are  $O(1)$  positions apart in  $T$ , resolving offset is  $O(1)$  time

# FM Index: problems solved

Solved! At the expense of adding some SA values ( $O(m)$  integers) to index  
Call this the "SA sample"

**(3)** Need a way to find where these occurrences are in  $T$ :

\$	a	b	a	a	b	<b>a<sub>0</sub></b>
<b>a<sub>0</sub></b>	\$	a	b	a	a	<b>b<sub>0</sub></b>
<b>a<sub>1</sub></b>	a	b	a	\$	a	<b>b<sub>1</sub></b>
<b>a<sub>2</sub></b>	b	a	\$	a	b	<b>a<sub>1</sub></b>
<b>a<sub>3</sub></b>	b	a	a	b	a	\$
<b>b<sub>0</sub></b>	a	\$	a	b	a	<b>a<sub>2</sub></b>
<b>b<sub>1</sub></b>	a	a	b	a	\$	<b>a<sub>3</sub></b>

**With SA sample we can do this in  
 $O(1)$  time per occurrence**



# FM Index: small memory footprint

Components of the FM Index:

First column ( $F$ ):	$\sim  \Sigma $ integers
Last column ( $L$ ):	$m$ characters
SA sample:	$m \cdot a$ integers, where $a$ is fraction of rows kept
Checkpoints:	$m \times  \Sigma  \cdot b$ integers, where $b$ is fraction of rows checkpointed

Example: DNA alphabet (2 bits per nucleotide),  $T$  = human genome,  $a = 1/32$ ,  $b = 1/128$

First column ( $F$ ):	16 bytes
Last column ( $L$ ):	2 bits * 3 billion chars = 750 MB
SA sample:	3 billion chars * 4 bytes/char / 32 = $\sim$ 400 MB
Checkpoints:	3 billion * 4 bytes/char / 128 = $\sim$ 100 MB
Total < 1.5 GB	