

Burrows-Wheeler Transform and FM Index

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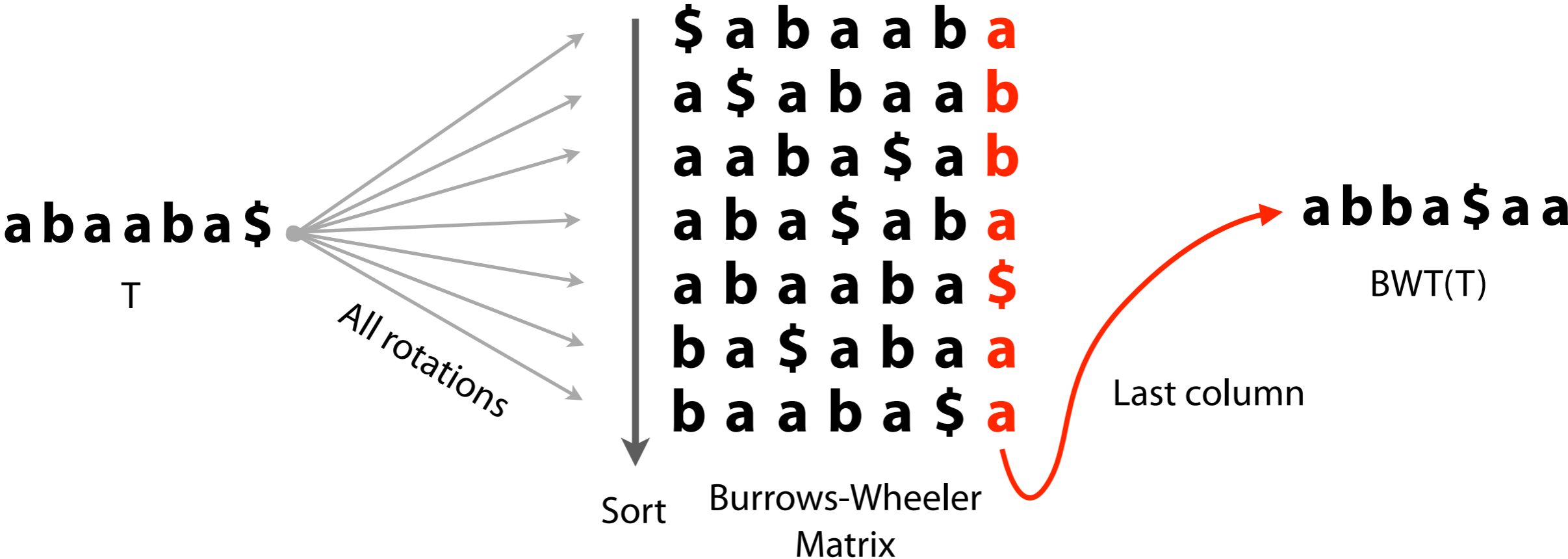
JOHNS HOPKINS

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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)) ]
```

Make list of all rotations

```
def bwm(t):  
    """ Return lexicographically sorted list of t's rotations """  
    return sorted(rotations(t))
```

Sort them

```
def bwtViaBwm(t):  
    """ Given T, returns BWT(T) by way of the BWM """  
    return ''.join(map(lambda x: x[-1], bwm(t)))
```

Take last column

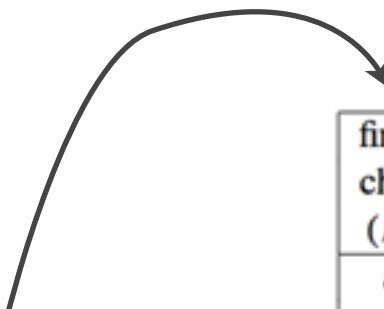
```
>>> bwtViaBwm("Tomorrow_and_tomorrow_and_tomorrow$")  
'w$wwdd__nnoooaattTmmmrrrrrrrooo__ooo'  
  
>>> bwtViaBwm("It_was_the_best_of_times_it_was_the_worst_of_times$")  
's$esttssffteww_hhmmbootttt_ii_woeearessIi_____  
  
>>> bwtViaBwm('in_the_jingle_jangle_morning_Ill_come_following_you$')  
'u_gleeeengj_mlh1_nnnnt$nwj__lggIolo_iiiiarfcmlylo_oo_'
```

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	\$
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)

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
```

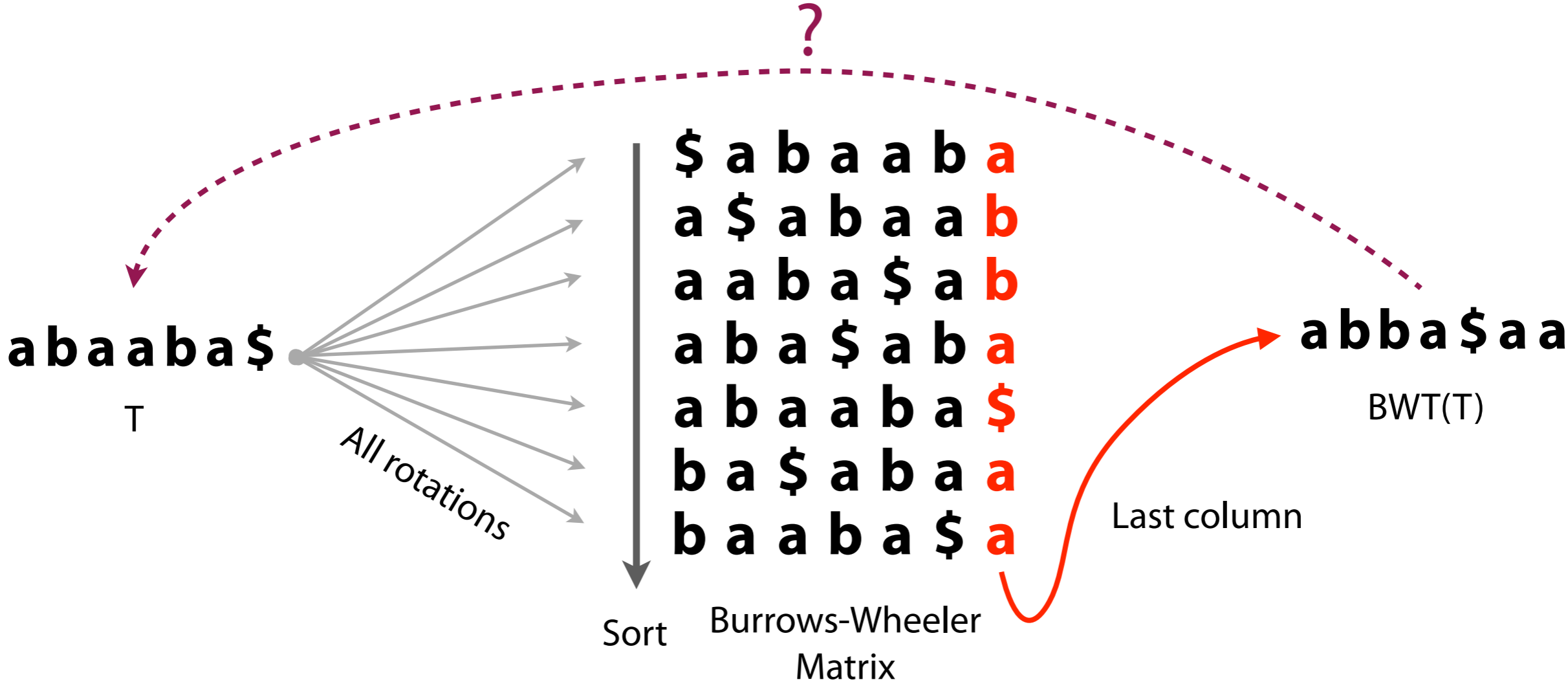
Take characters just
to the left of the
sorted suffixes

```
>>> bwtViaSa("Tomorrow_and_tomorrow_and_tomorrow$")  
'w$wwdd__nnoooaattTmmrrrrrrrooo__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.

a₀ **b**₀ **a**₁ **a**₂ **b**₁ **a**₃ \$

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

Burrows-Wheeler Transform

BWM with T-ranking:

	<i>F</i>						<i>L</i>
	\$	a ₀	b ₀	a ₁	a ₂	b ₁	a₃
	a₃	\$	a ₀	b ₀	a ₁	a ₂	b ₁
	a₁	a ₂	b ₁	a ₃	\$	a ₀	b ₀
	a₂	b ₁	a ₃	\$	a ₀	b ₀	a₁
	a₀	b ₀	a ₁	a ₂	b ₁	a ₃	\$
	b ₁	a ₃	\$	a ₀	b ₀	a ₁	a₂
	b ₀	a ₁	a ₂	b ₁	a ₃	\$	a₀

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

And look at just the **a**s

as occur in the same order in *F* and *L*. As we look down columns, in both cases we see: **a₃, a₁, a₂, a₀**

Burrows-Wheeler Transform

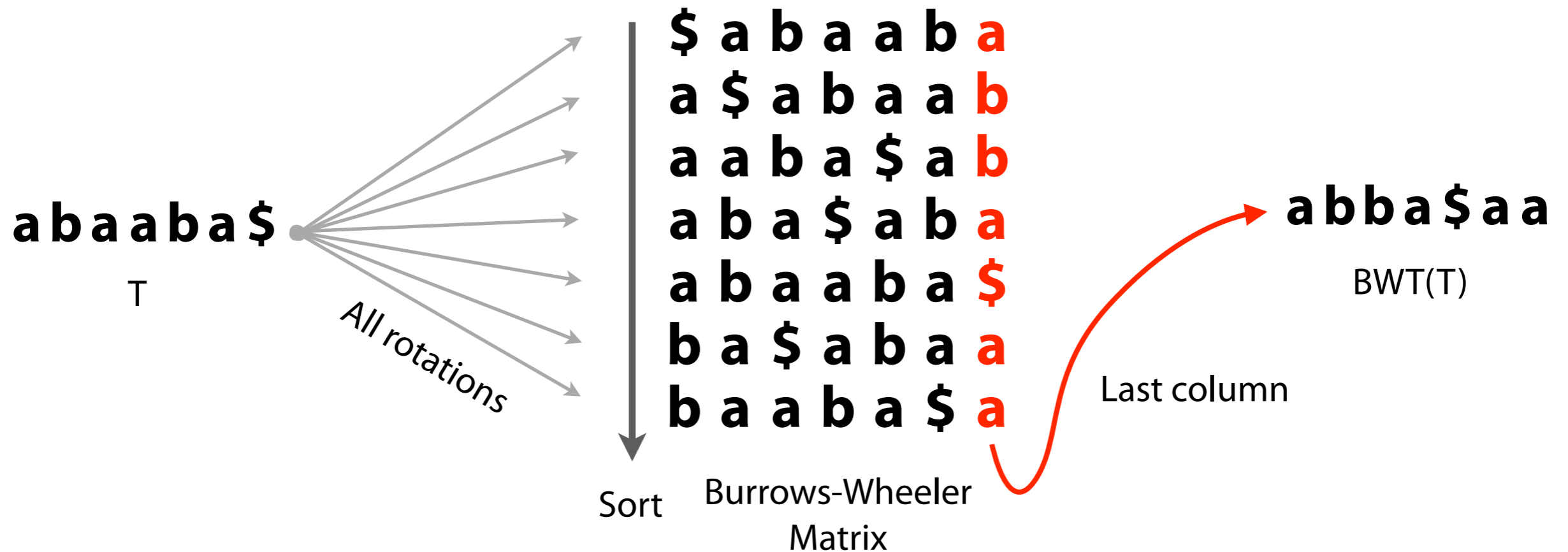
BWM with T-ranking:

<i>F</i>						<i>L</i>
\$	a ₀	b ₀	a ₁	a ₂	b ₁	a ₃
a ₃	\$	a ₀	b ₀	a ₁	a ₂	b₁
a ₁	a ₂	b ₁	a ₃	\$	a ₀	b₀
a ₂	b ₁	a ₃	\$	a ₀	b ₀	a ₁
a ₀	b ₀	a ₁	a ₂	b ₁	a ₃	\$
b₁	a ₃	\$	a ₀	b ₀	a ₁	a ₂
b₀	a ₁	a ₂	b ₁	a ₃	\$	a ₀

Same with **b**s: **b₁**, **b₀**

Burrows-Wheeler Transform

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



How is it useful for compression?

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How is it an index?

Burrows-Wheeler Transform: LF Mapping

BWM with T-ranking:

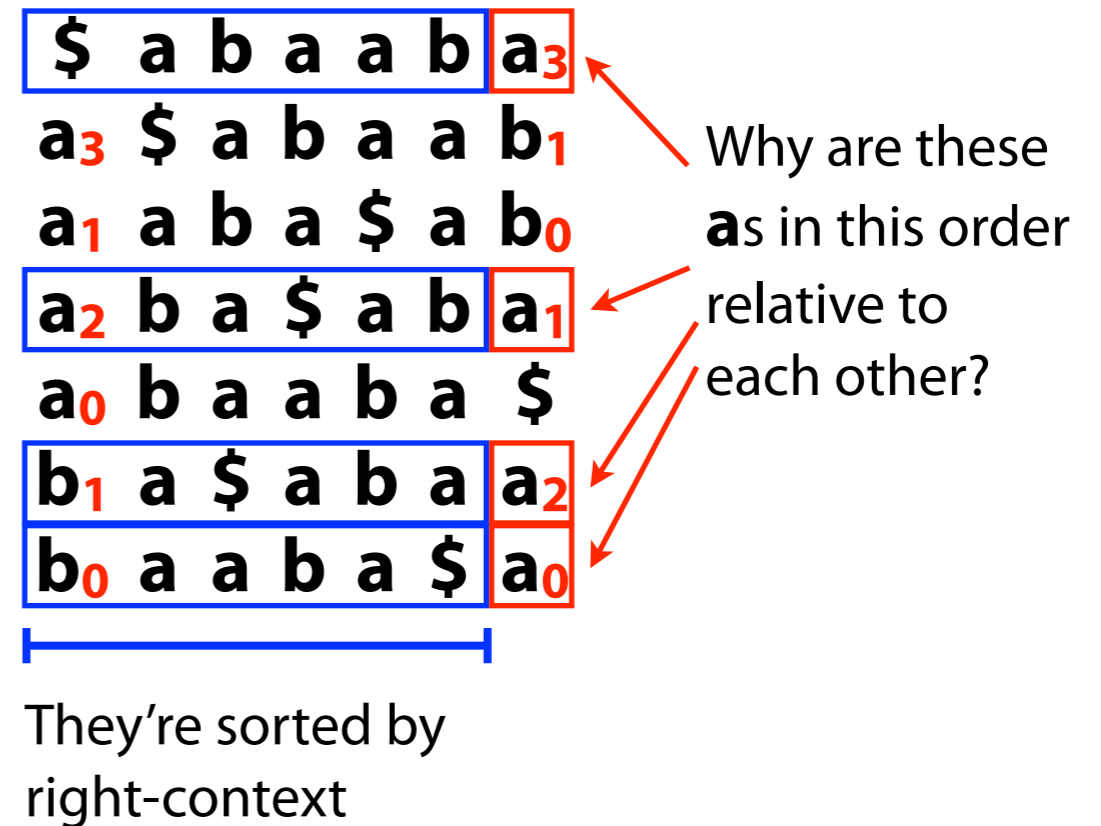
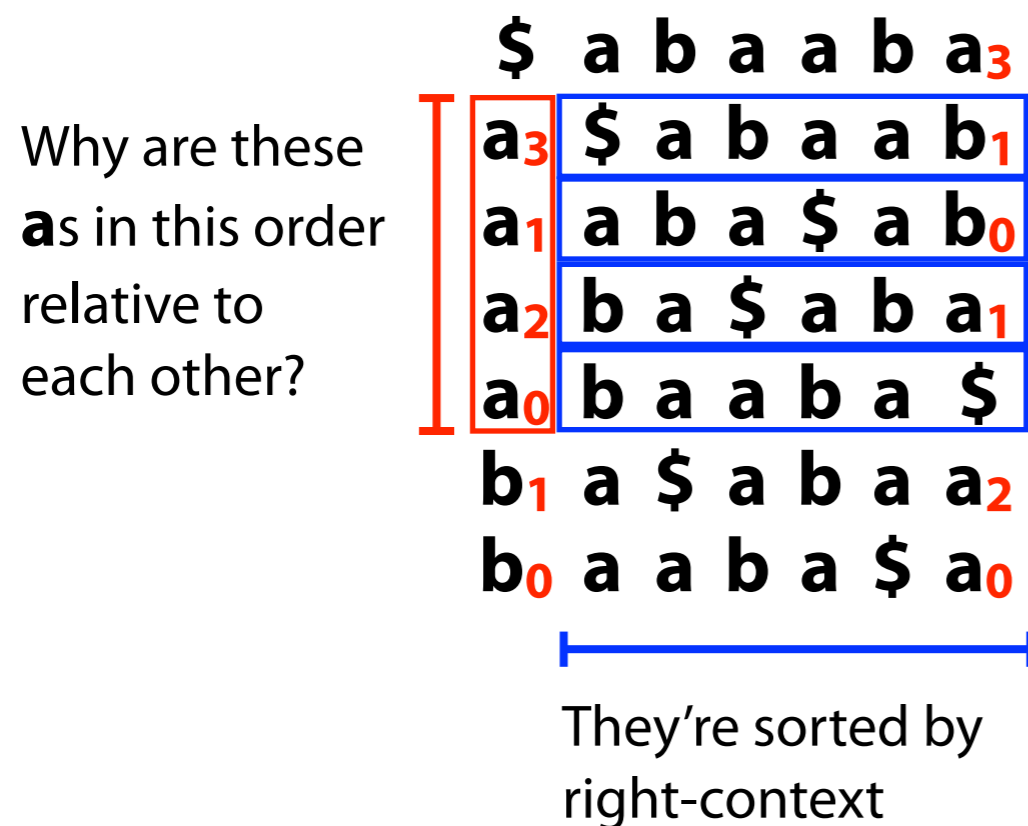
F						L
\$	a₀	b₀	a₁	a₂	b₁	a₃
a₃	\$	a₀	b₀	a₁	a₂	b₁
a₁	a₂	b₁	a₃	\$	a₀	b₀
a₂	b₁	a₃	\$	a₀	b₀	a₁
a₀	b₀	a₁	a₂	b₁	a₃	\$
b₁	a₃	\$	a₀	b₀	a₁	a₂
b₀	a₁	a₂	b₁	a₃	\$	a₀

LF Mapping: The i^{th} occurrence of a character c in L and the i^{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

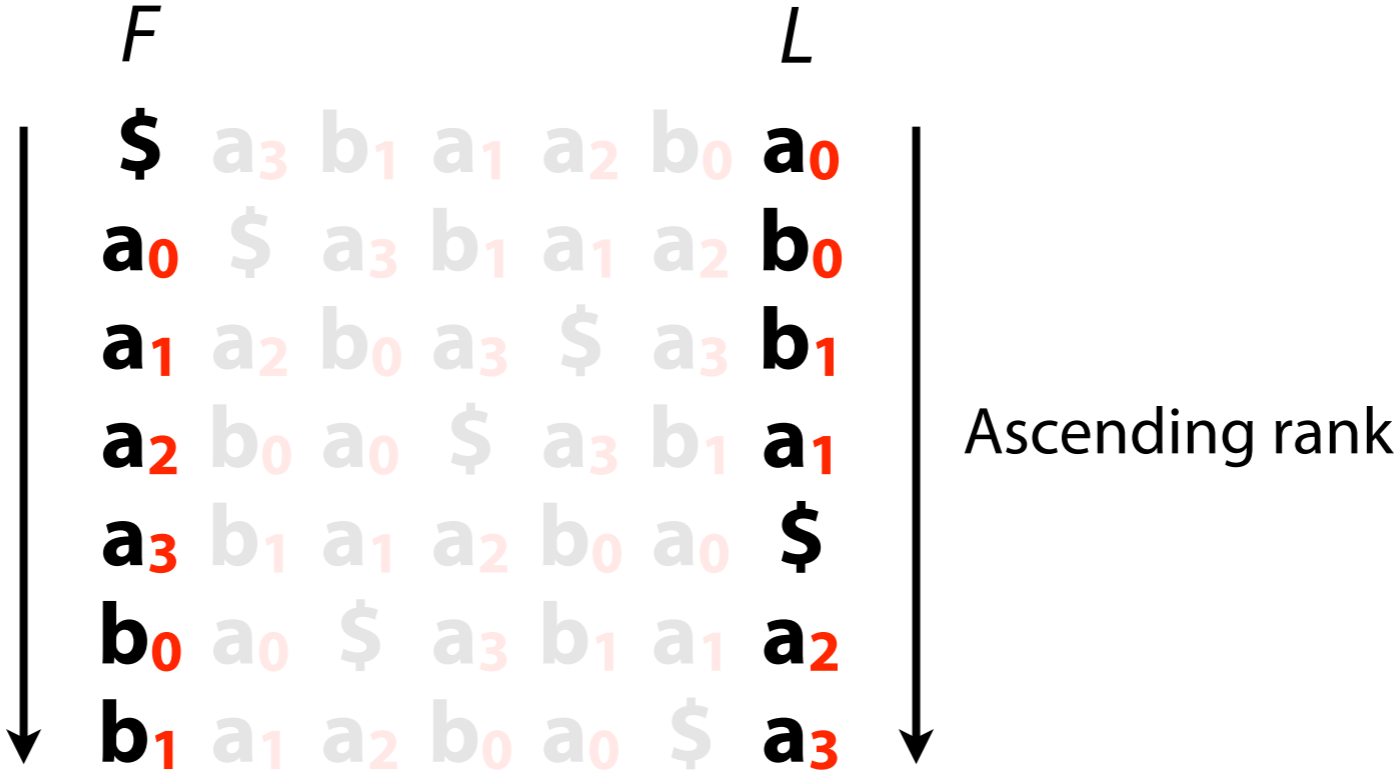
BWM with T-ranking:

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

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 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

F *L*

\$ **a₀**

a₀ **b₀**

a₁ **b₁** ← Which BWM row *begins* with **b₁**?

a₂ **a₁** Skip row starting with **\$** (1 row)

a₃ **\$** Skip rows starting with **a** (4 rows)

Skip row starting with **b₀** (1 row)

b₀ **a₂**

Answer: row 6

row 6 → **b₁** **a₃**

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**₁₀₀? (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 $\$$: $\mathbf{a_0}$

$\mathbf{a_0}$: LF Mapping says this is same occurrence of \mathbf{a} as first \mathbf{a} in F . **Jump** to row *beginning* with $\mathbf{a_0}$. L contains character just **prior** to $\mathbf{a_0}$: $\mathbf{b_0}$.

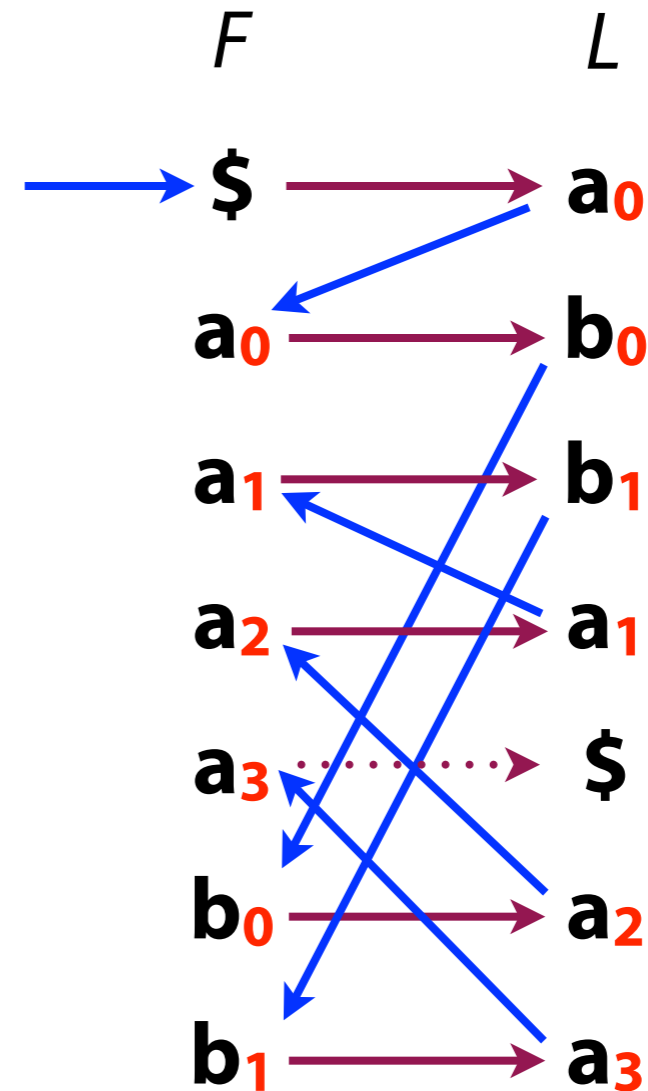
Repeat for $\mathbf{b_0}$, get $\mathbf{a_2}$

Repeat for $\mathbf{a_2}$, get $\mathbf{a_1}$

Repeat for $\mathbf{a_1}$, get $\mathbf{b_1}$

Repeat for $\mathbf{b_1}$, get $\mathbf{a_3}$

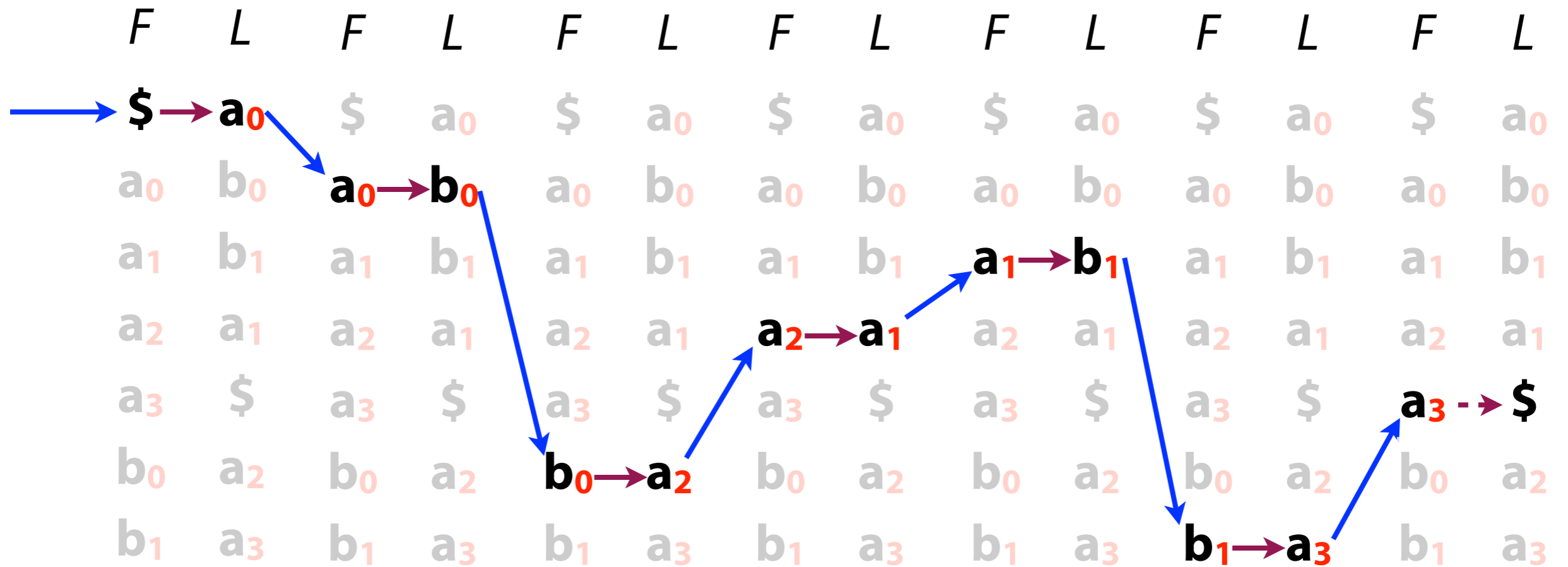
Repeat for $\mathbf{a_3}$, get $\mathbf{\$}$, done



Reverse of chars we visited = $\mathbf{a_3 b_1 a_1 a_2 b_0 a_0 \$} = T$

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$wwdd__nnoooaattTmmrrrrrrrooo__ooo")
'Tomorrow_and_tomorrow_and_tomorrow$'

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

>>> 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 auxiliary data structures*

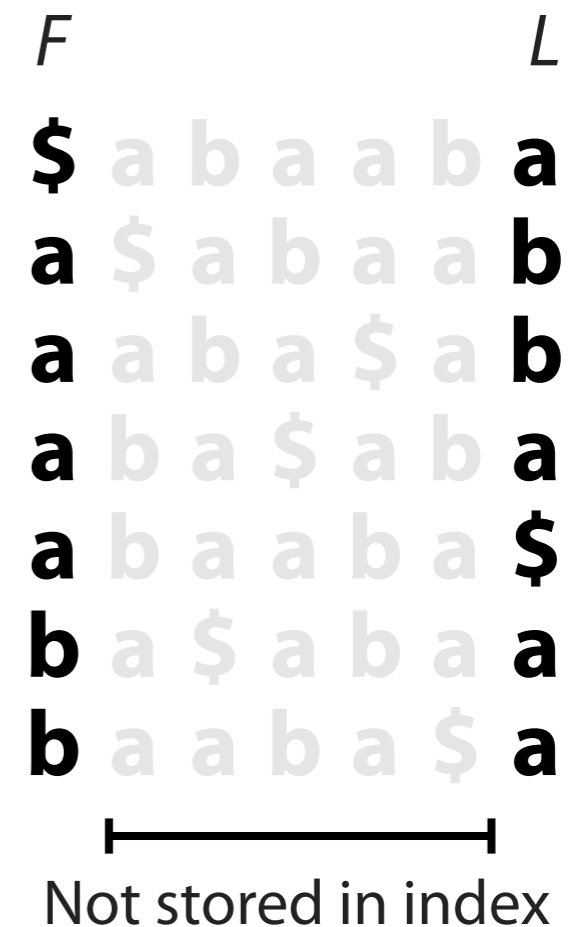
“FM” supposedly stands for “Full-text Minute-space.”
(But inventors are named Ferragina and Manzini)

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!

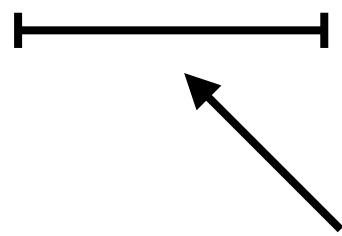


Paolo Ferragina, and Giovanni Manzini. "Opportunistic data structures with applications." *Foundations of Computer Science, 2000. Proceedings. 41st Annual Symposium on.* IEEE, 2000.

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

$P = \mathbf{aba}$

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**

$P = \mathbf{ab}a$

$P = a\mathbf{ba}$

<i>F</i>						<i>L</i>
\$	a	b	a	a	b	a₀
a₀	\$	a	b	a	a	b₀
a₁	a	b	a	\$	a	b₁
a₂	b	a	\$	a	b	a₁
a₃	b	a	a	b	a	\$
b₀	a	\$	a	b	a	a₂
b₁	a	a	b	a	\$	a₃

← Look at those rows in *L*.
b₀, **b₁** are **b**s occurring just to left.

Use LF Mapping. Let new range delimit those **b**s

b₀	a	\$	a	b	a	a₂
b₁	a	a	b	a	\$	a₃

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

FM Index: querying

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

$P = \mathbf{aba}$

$P = \mathbf{aba}$

<i>F</i>						<i>L</i>
\$	a	b	a	a	b	a₀
a₀	\$	a	b	a	a	b₀
a₁	a	b	a	\$	a	b₁
a₂	b	a	\$	a	b	a₁
a₃	b	a	a	b	a	\$
b₀	a	\$	a	b	a	a₂
b₁	a	a	b	a	\$	a₃

← **a₂**, **a₃** occur just to left.

Use LF Mapping →

<i>F</i>						<i>L</i>
\$	a	b	a	a	b	a₀
a₀	\$	a	b	a	a	b₀
a₁	a	b	a	\$	a	b₁
a₂	b	a	\$	a	b	a₁
a₃	b	a	a	b	a	\$
b₀	a	\$	a	b	a	a₂
b₁	a	a	b	a	\$	a₃

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

FM Index: querying

$P = \mathbf{aba}$

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

	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
$[3, 5)$	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

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 \$

Where are these?

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 $\mathbf{a_0}$
	$\mathbf{a_0}$	\$	a	b	a	a $\mathbf{b_0}$
	$\mathbf{a_1}$	a	b	a	\$	a $\mathbf{b_1}$
	$\mathbf{a_2}$	b	a	\$	a	b $\mathbf{a_1}$
	$\mathbf{a_3}$	b	a	a	b	a \$
Rows with \mathbf{ba} prefix	$\mathbf{b_0}$	a	\$	a	b	a $\mathbf{a_2}$
	$\mathbf{b_1}$	a	a	b	a	\$ $\mathbf{a_3}$

← No \mathbf{bs} !

FM Index: querying

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

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

<i>F</i>						<i>L</i>
\$	a	b	a	a	b	a₃
a₀	\$	a	b	a	a	b₁
a₁	a	b	a	\$	a	b₀
a₂	b	a	\$	a	b	a₁
a₃	b	a	a	b	a	\$
b₀	a	\$	a	b	a	a₂
b₁	a	a	b	a	\$	a₀

Scan, looking for **b**s

FM Index: lingering issues

(1) Scanning for preceding character is slow

	\$	a	b	a	a	b	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	\$	\$
b ₀	a	\$	a	b	a	a	a ₂
b ₁	a	a	b	a	\$	a	a ₃

$O(m)$
 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	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	\$	\$
b ₀	a	\$	a	b	a	a	a ₂
b ₁	a	a	b	a	\$	a	a ₃

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	a₀
	a₀	\$ a b a a	b₀
	a₁	a b a \$ a	b₁
	a₂	b a \$ a b	a₁
	a₃	b a a b a	\$
	b₀	a \$ a b a	a₂
	b₁	a a b a \$	a₃

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

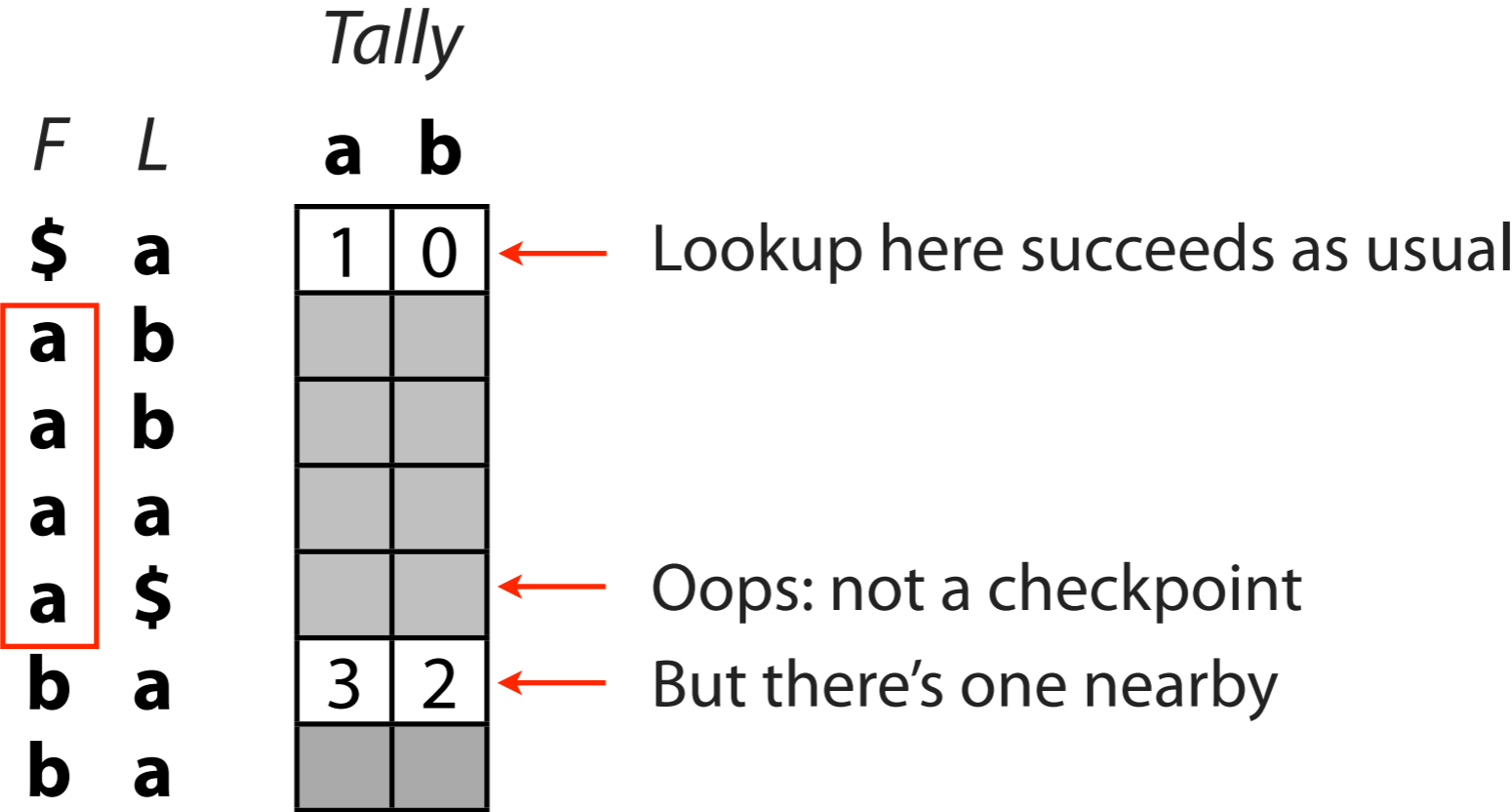
	<i>F</i>	<i>L</i>	<i>Tally</i>	
	\$	a	a	b
	a	b	1	0
	a	b	1	1
	a	a	1	2
	a	a	2	2
	a	\$	2	2
	b	a	3	2
	b	a	4	2

We infer **b₀** and **b₁** appear in *L* in this range

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

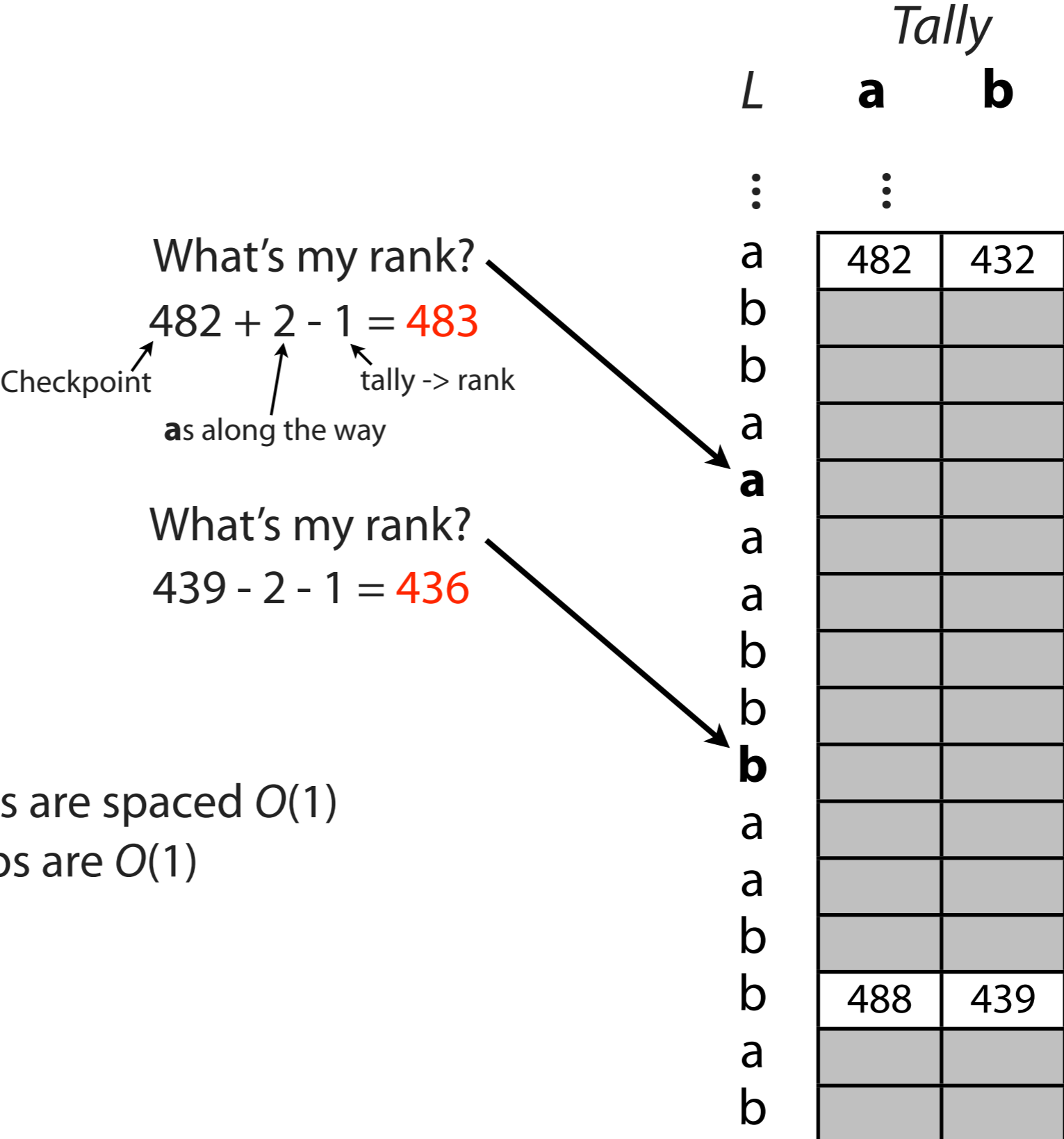
FM Index: fast rank calculations

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



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



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)

<i>F</i>		<i>L</i>	
\$	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

```
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

With checkpoints, we greatly reduce # integers needed for ranks

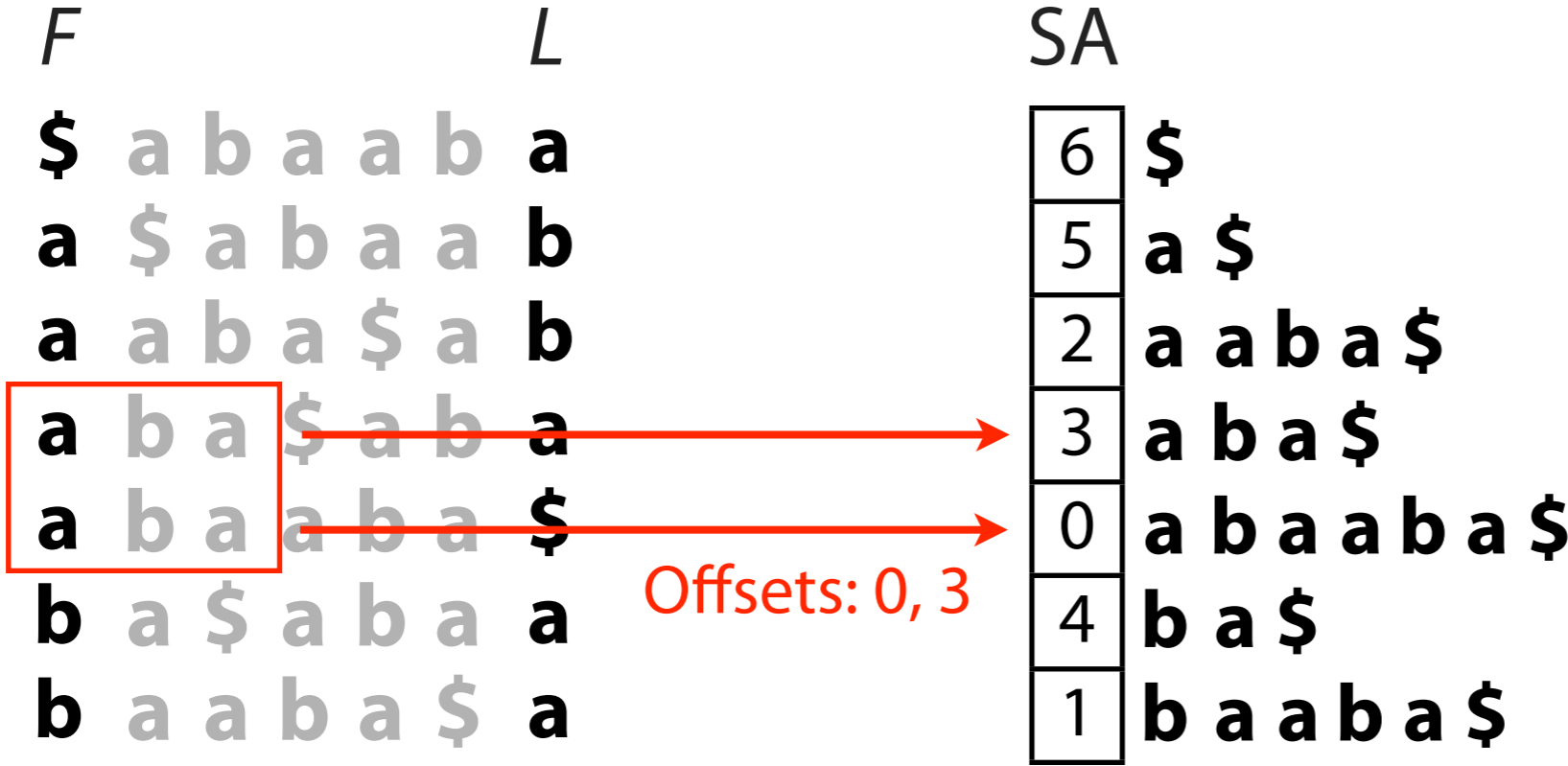
But it's still $O(m)$ space - there's literature on how to improve this space bound

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 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

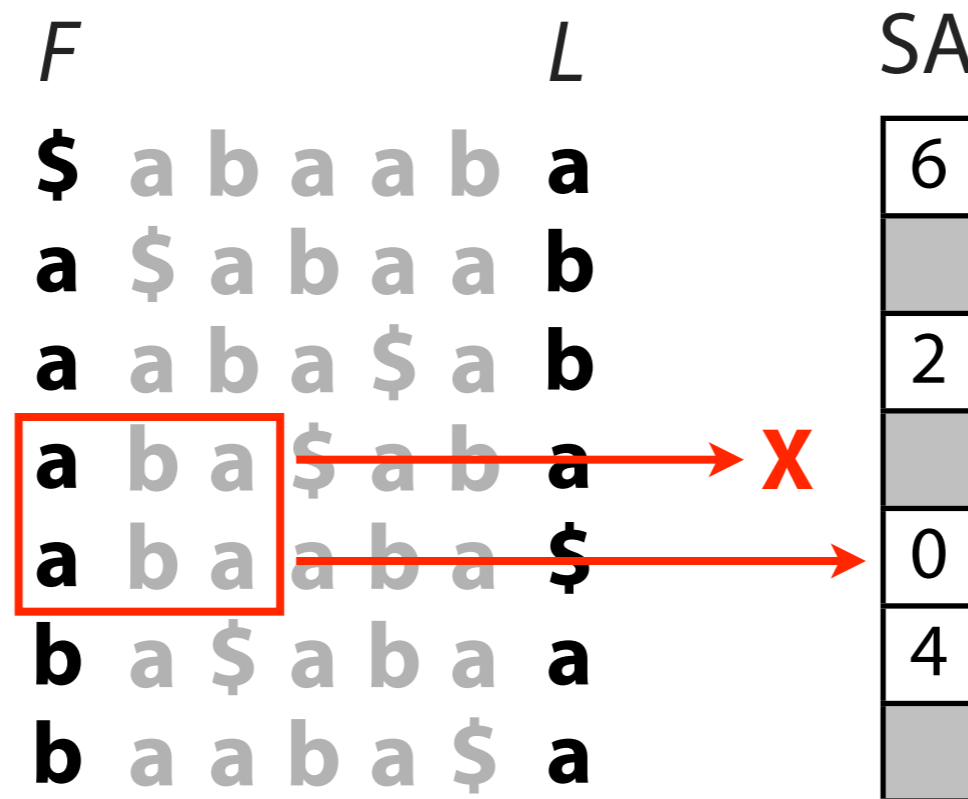
If suffix array were part of index, we could simply look up the offsets



But SA requires m integers

FM Index: resolving offsets

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

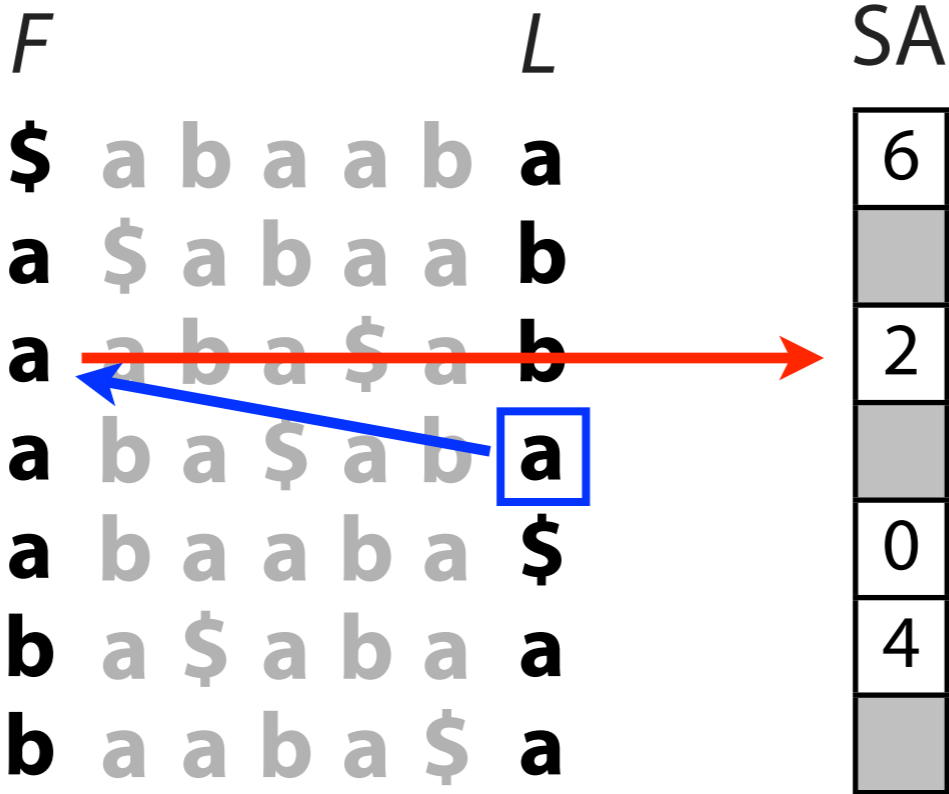


Lookup for row 4 succeeds - we kept that entry of SA

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



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	a₀
a₀	\$	a	b	a	a	b₀
a₁	a	b	a	\$	a	b₁
a₂	b	a	\$	a	b	a₁
a₃	b	a	a	b	a	\$
b₀	a	\$	a	b	a	a₂
b₁	a	a	b	a	\$	a₃

**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