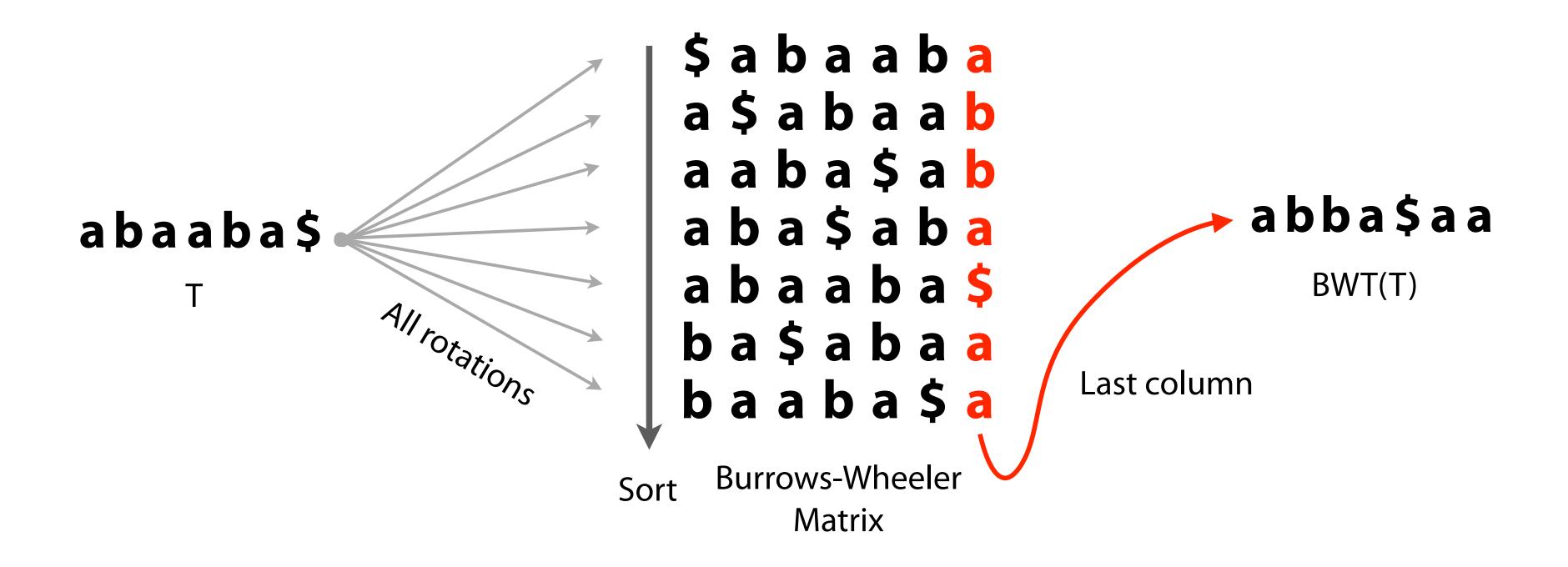
BWT & FM INDEX



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?

```
def rotations(t):
   """ Return list of rotations of input string t """
                                                            Make list of all rotations
   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
                                                            Sort them
   return sorted(rotations(t))
def bwtViaBwm(t):
   """ Given T, returns BWT(T) by way of the BWM """
                                                            Take last column
   return ''.join(map(lambda x: x[-1], bwm(t)))
>>> bwtViaBwm("Tomorrow_and_tomorrow_and_tomorrow$")
 'w$wwdd nnoooaattTmmmrrrrrooo ooo'
>>> bwtViaBwm("It_was_the_best_of_times_it_was_the_worst_of_times$")
 's$esttssfftteww hhmmbootttt ii woeeaaressIi
>>> bwtViaBwm('in_the_jingle_jangle_morning_Ill_come_following_you$')
 'u gleeeengj mlhl nnnnt$nwj lggIolo iiiiarfcmylo oo '
```

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	sorted rotations
(<i>L</i>)	
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
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
0	n with Huffman or arithmetic coding. Bri
0	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.

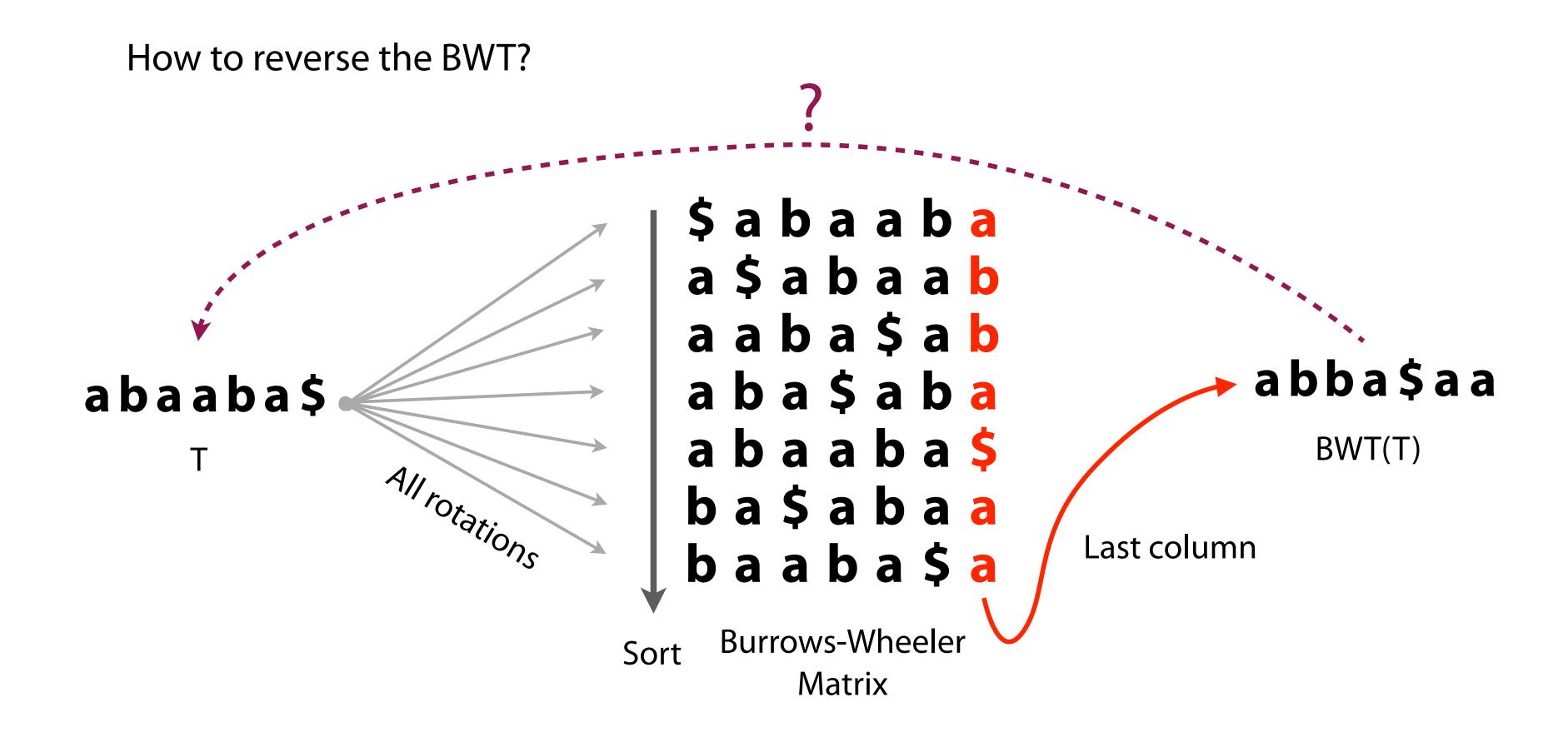
BWM bears a resemblance to the suffix array

Sort order is the same whether rows are rotations or suffixes

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"



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 b₀ a₁ a₂ b₁ a₃ \$

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

Note: we *do not* actually write this information in the text / BWM, we Are simply including it here to help us track "which" occurrences of each character in the BWM correspond to the occurrences in the text.

```
F L

BWM with T-ranking: $ 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

b_0 a_1 a_2 b_1 a_3 $ a_0

b_0 a_1 a_2 b_1 a_3 $ a_0
```

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: \mathbf{a}_3 , \mathbf{a}_1 , \mathbf{a}_2 , \mathbf{a}_0

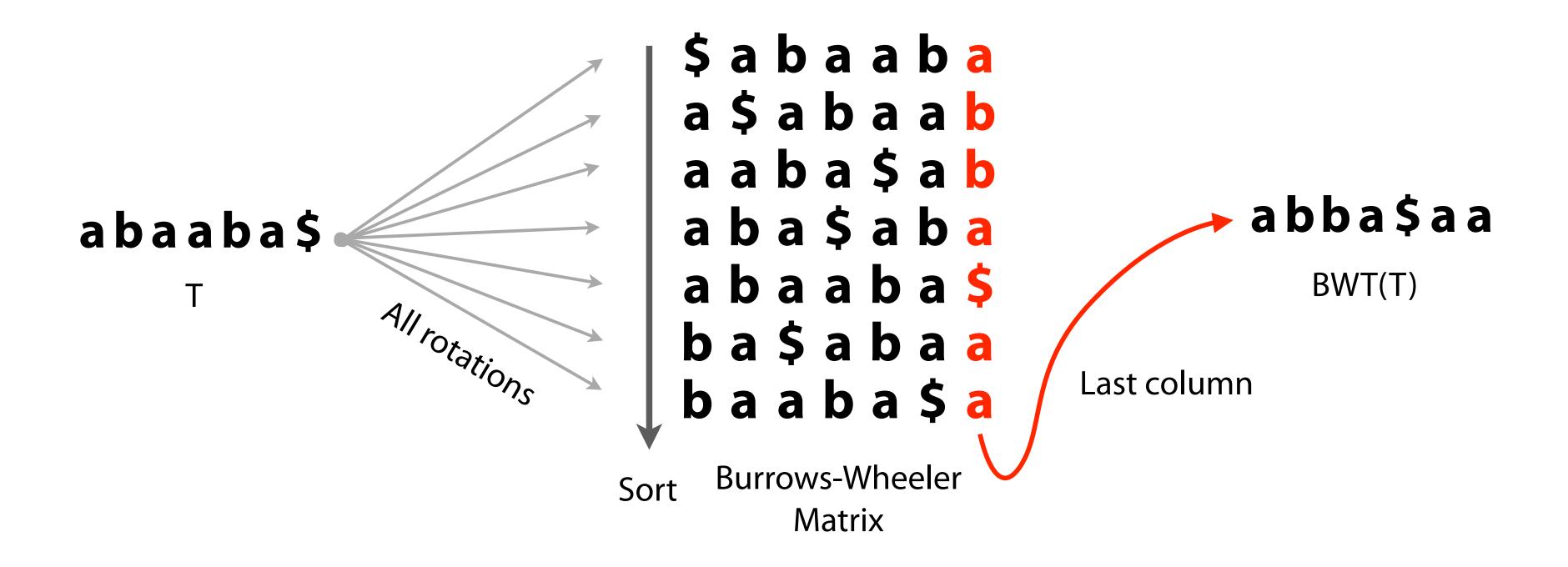
```
F

BWM with T-ranking:

$ 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_0 b_0 a_1 a_2 b_0 a_0 b_0 a_1 a_2 b_0 a_0 b_0 a_1 a_2 b_0
```

Same with **b**s: **b**₁, **b**₀

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



How is it useful for compression?

How is it reversible?

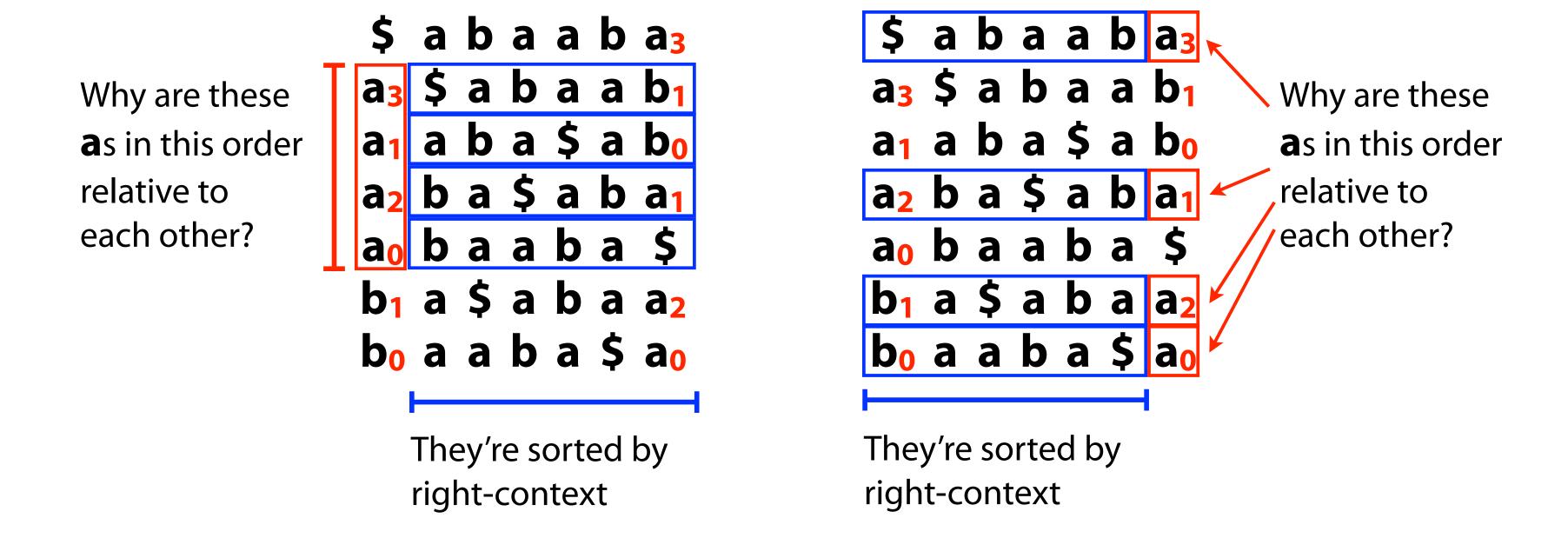
How is it an index?

```
BWM with T-ranking: $ 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_3 $ a_0 b_0 a_1 a_2 b_0 a_1 a_2 b_1 a_3 $ a_0 b_0 a_1 a_2 b_0 a_1 a_2 b_1 a_3 $ a_0 b_0 a_1 a_2 b_0 a_1 a_2 b_1 a_3 $ a_0 b_0 a_1 a_2 b_0 a_1 a_2 b_1 a_3 $ a_0 b_0 a_1 a_2 b_0 a_1 a_2 b_1 a_3 $ a_0 b_0 a_1 a_2 b_0 a_1 a_2 b_1 a_3 $ a_0 b_0 a_1 a_2 b_0 a_1 a_2 b_1 a_3 $ a_0 b_0 a_1 a_2 b_0 a_1 a_2 b_1 a_3 $ a_0 b_0 a_1 a_2 b_0 a_1 a_2 b_1 a_3 $ a_0 b_0 a_1 a_2 b_0
```

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

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

BWM with 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>
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...

BWM with B-ranking:

```
$ 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

Ascending rank

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>3</sub> b<sub>1</sub> a<sub>1</sub> a<sub>2</sub>
```

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

```
a<sub>0</sub>
           b<sub>0</sub>
a<sub>0</sub>
                          Which BWM row begins with b<sub>1</sub>?
a<sub>1</sub>
                              Skip row starting with $ (1 row)
           a<sub>1</sub>
a<sub>2</sub>
                              Skip rows starting with a (4 rows)
a<sub>3</sub>
                              Skip row starting with b_0 (1 row)
bo
           a<sub>2</sub>
                              Answer: row 6
```

Say *T* has 300 **A**s, 400 **C**s, 250 **G**s and 700 **T**s and \$ < **A** < **C** < **G** < **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 \mathbf{G} (100 rows)

Answer: row 1 + 300 + 400 + 100 = 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 \$: ao

a₀: LF Mapping says this is same occurrence of **a** as first **a** in *F*. Jump to row *beginning* with **a**₀. *L* contains character just prior to **a**₀: **b**₀.

Repeat for **b**₀, get **a**₂

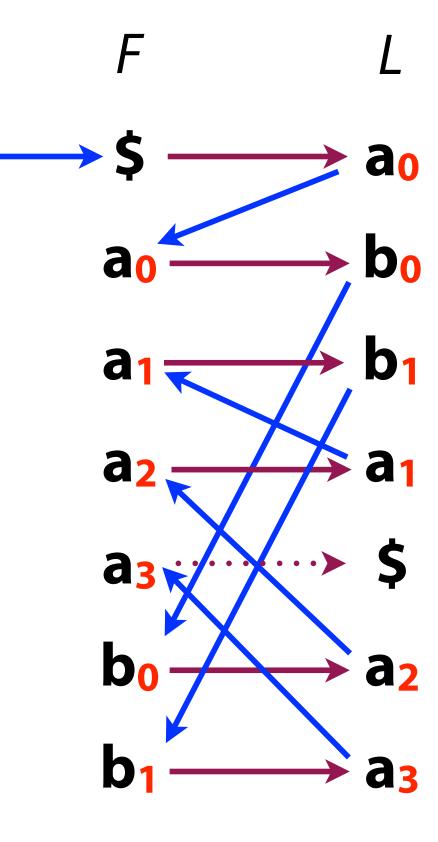
Repeat for a2, get a1

Repeat for **a**₁, get **b**₁

Repeat for **b**₁, get **a**₃

Repeat for **a**₃, get \$, done

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



Burrows-Wheeler Transform: reversing

Another way to visualize reversing BWT(T):

Burrows-Wheeler Transform: reversing

```
>>> reverseBwt("w$wwdd__nnoooaattTmmmrrrrrooo__ooo")
'Tomorrow_and_tomorrows'
>>> reverseBwt("s$esttssfftteww_hhmmbootttt_ii__woeeaaressIi_____")
'It_was_the_best_of_times_it_was_the_worst_of_times$'
>>> reverseBwt("u_gleeeengj_mlhl_nnnnt$nwj__lggIolo_iiiiarfcmylo_oo_")
'in_the_jingle_jangle_morning_Ill_come_following_you$'
```

```
def reverseBwt(bw):
    ''' Make T from BWT(T) '''
    ranks list is m integers
long! We'll fix later.

promote the start of the s
```

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)

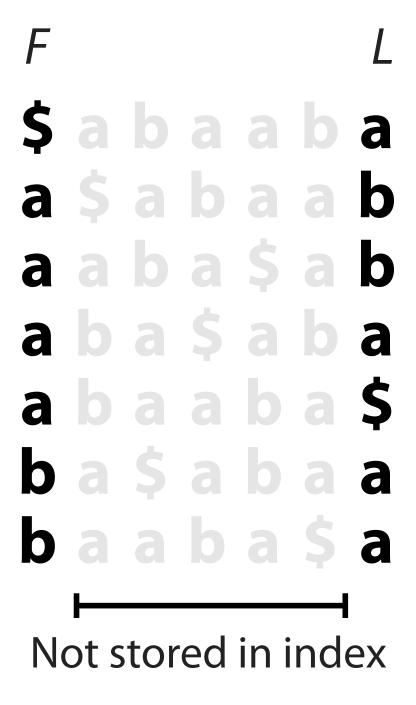
Core of index consists of *F* and *L* from BWM:

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

And *L* is compressible

Potentially very space-economical!

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



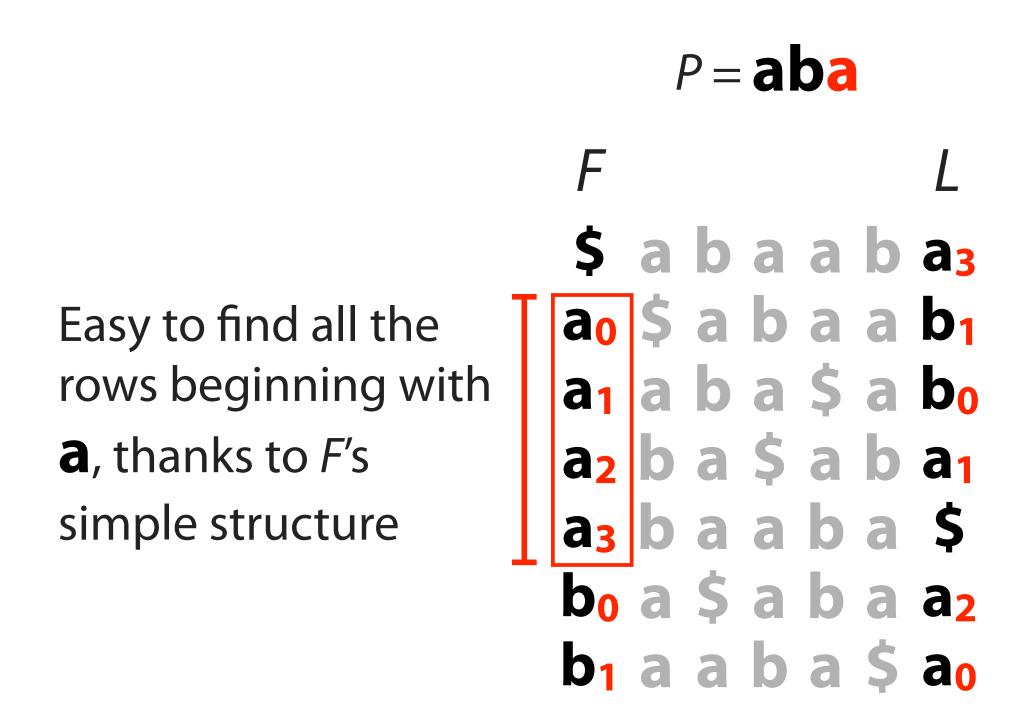
Though BWM is related to suffix array, we can't query it the same way

```
$ a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b a a b
```

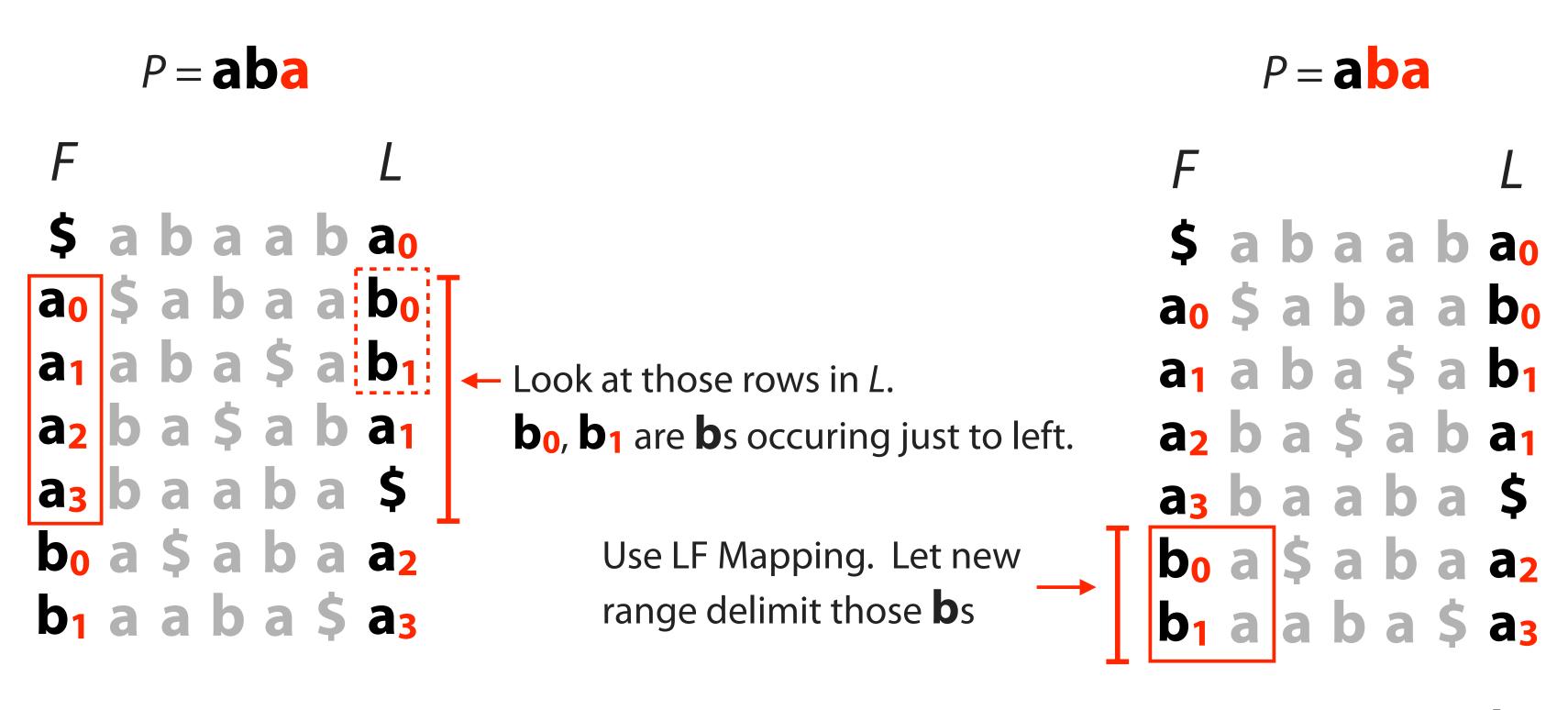
We don't have these columns; binary search isn't possible

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

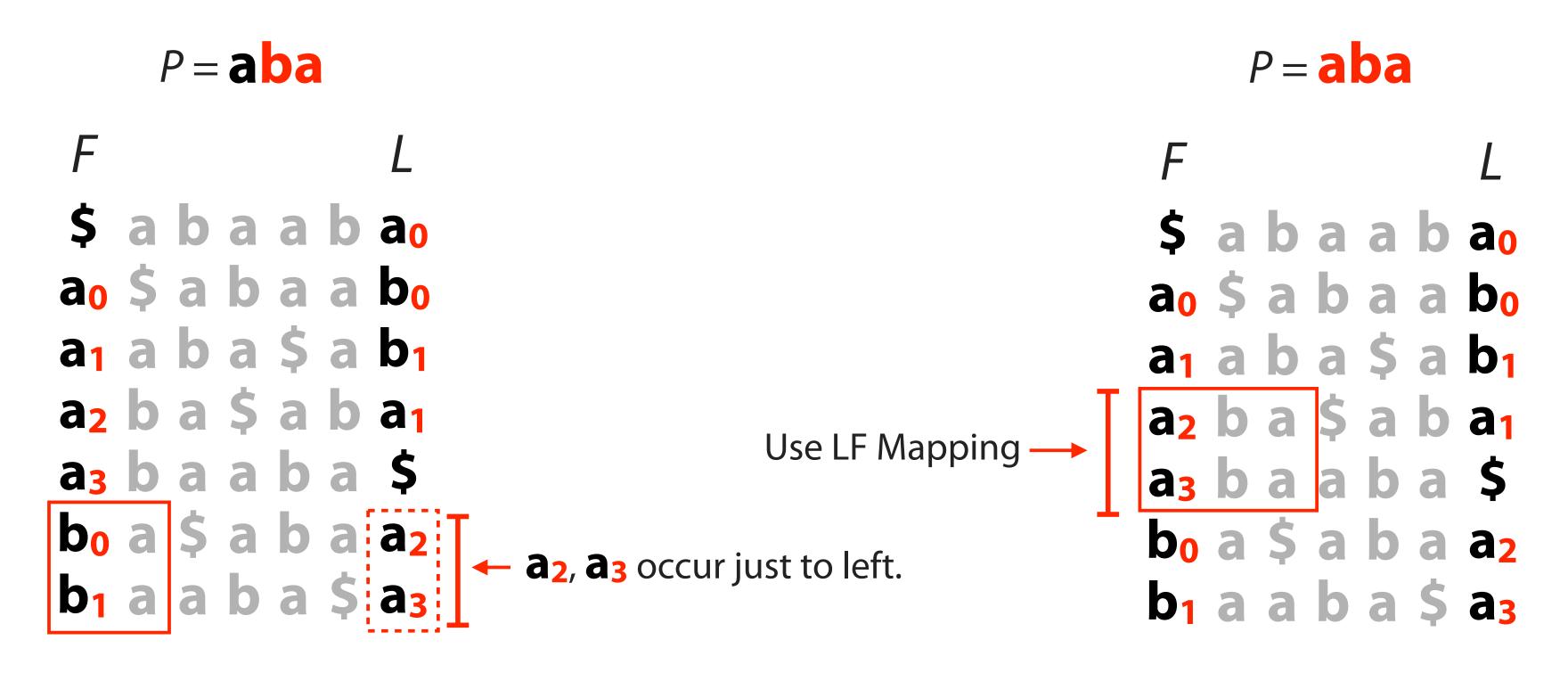


We have rows beginning with **a**, now we seek rows beginning with **ba**



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

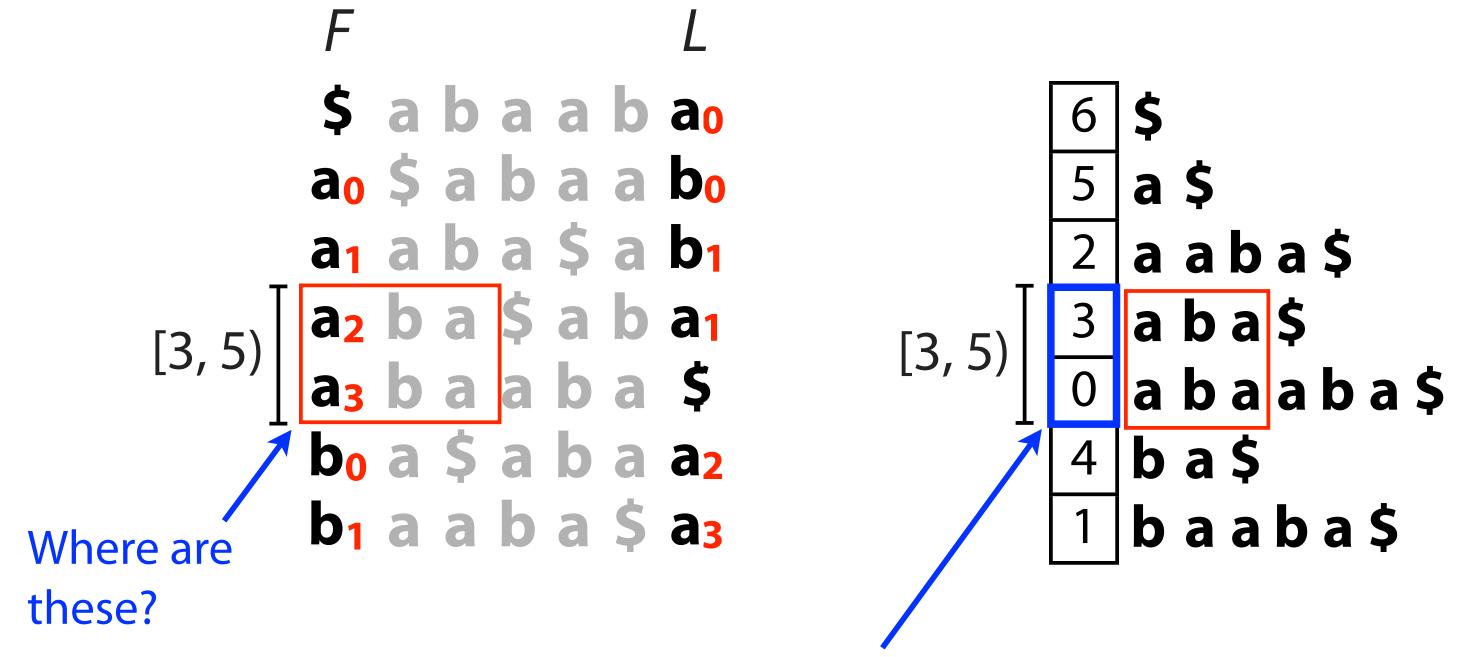
We have rows beginning with **ba**, now we seek rows beginning with **aba**



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

P = aba

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



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

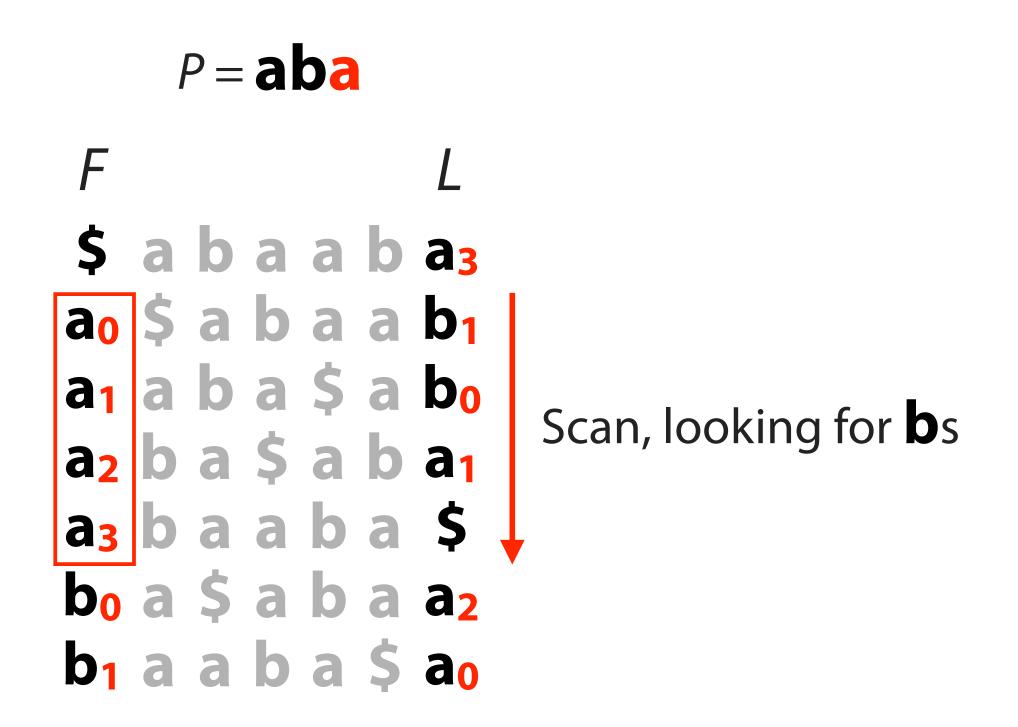
When *P* does not occur in *T*, we will eventually fail to find the next character in *L*:

$$P = \mathbf{bba}$$

$$F \qquad \qquad L$$

$$\mathbf{\$} \ \mathbf{a} \ \mathbf{b} \ \mathbf{a} \ \mathbf{a} \ \mathbf{b} \ \mathbf{a} \ \mathbf{a} \ \mathbf{b} \ \mathbf{a} \ \mathbf{b} \ \mathbf{a} \ \mathbf{a} \ \mathbf{b} \ \mathbf{a}$$

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



FM Index: lingering issues

(1) Scanning for preceding character is slow



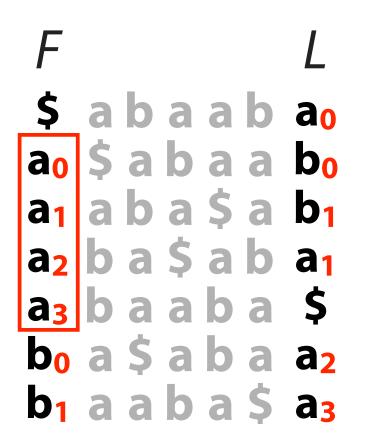
(2) Storing ranks takes too much space

```
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
```

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

```
$ abaab a<sub>0</sub>
a<sub>0</sub> $ abaab a<sub>0</sub>
a<sub>1</sub> abaab a<sub>5</sub>
a<sub>1</sub> abaab a<sub>1</sub>
a<sub>2</sub> baabaab a<sub>1</sub>
a<sub>3</sub> baabaa$
b<sub>0</sub> a$ abaa a<sub>2</sub>
b<sub>1</sub> aaba$ a<sub>3</sub>
```

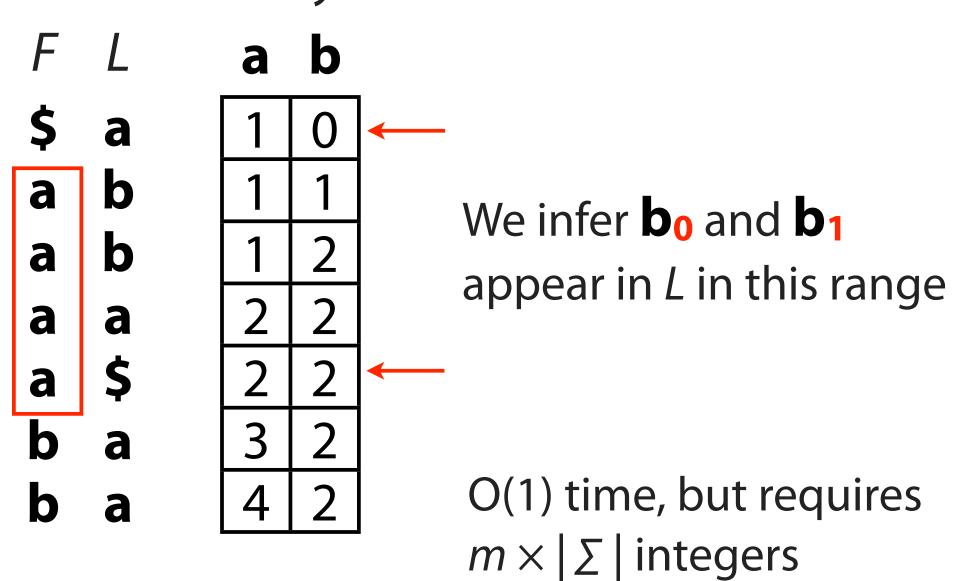
Is there an O(1) way to determine which **b**s precede the **a**s in our range?



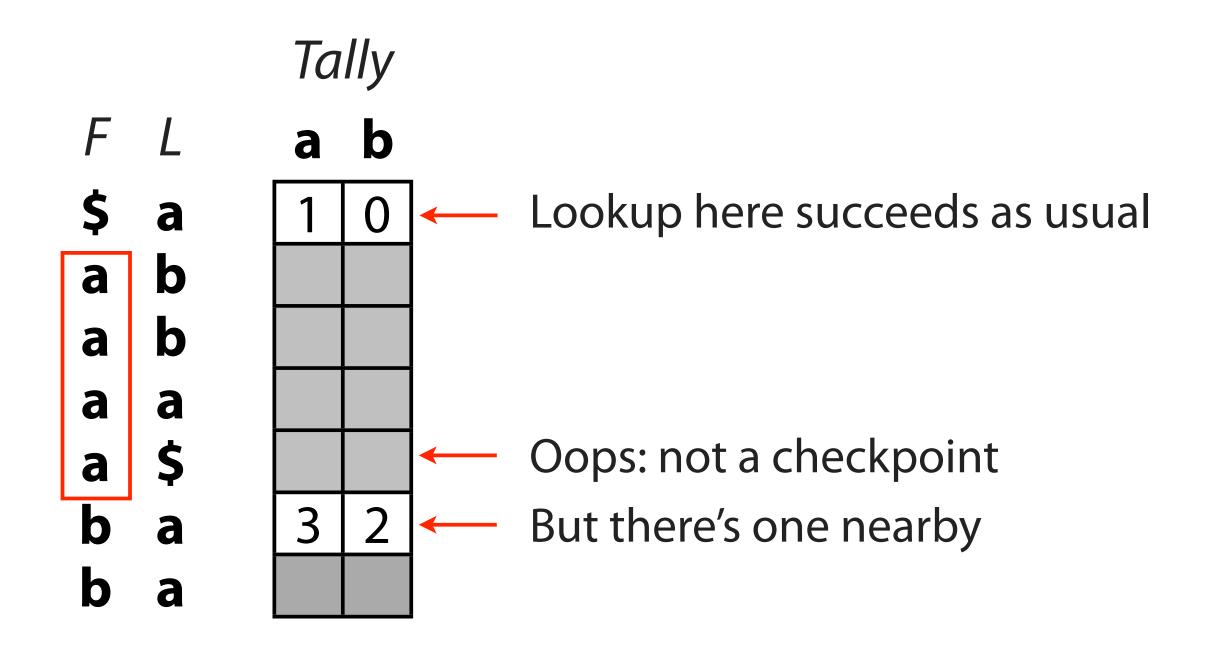
Occ(c, k) = # of times c occurs in the first k characters of BWT(S), aka the LF mapping.

Tally — also referred to as Occ(c, k)

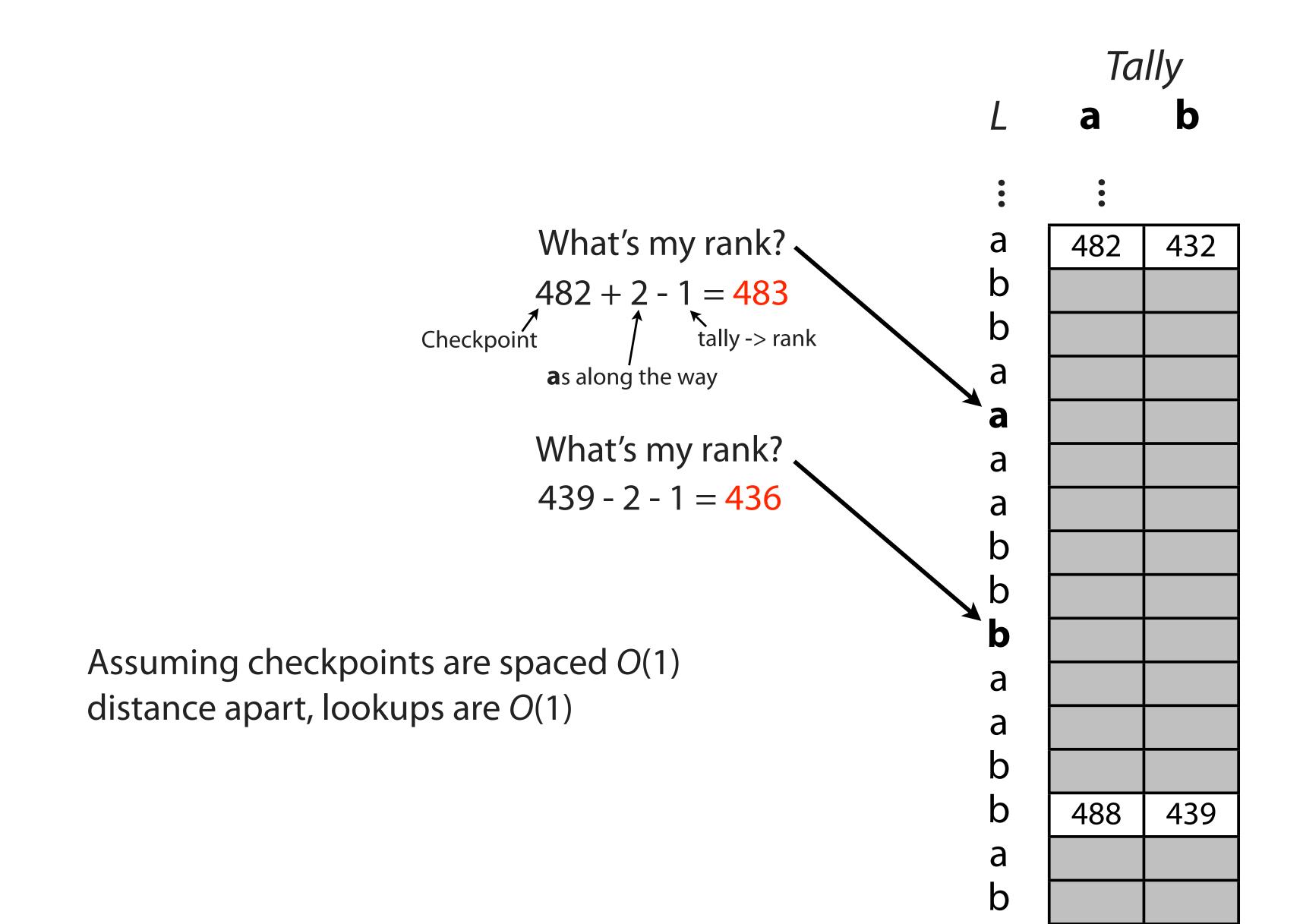
Idea: pre-calculate # **a**s, **b**s in *L* up to every row:



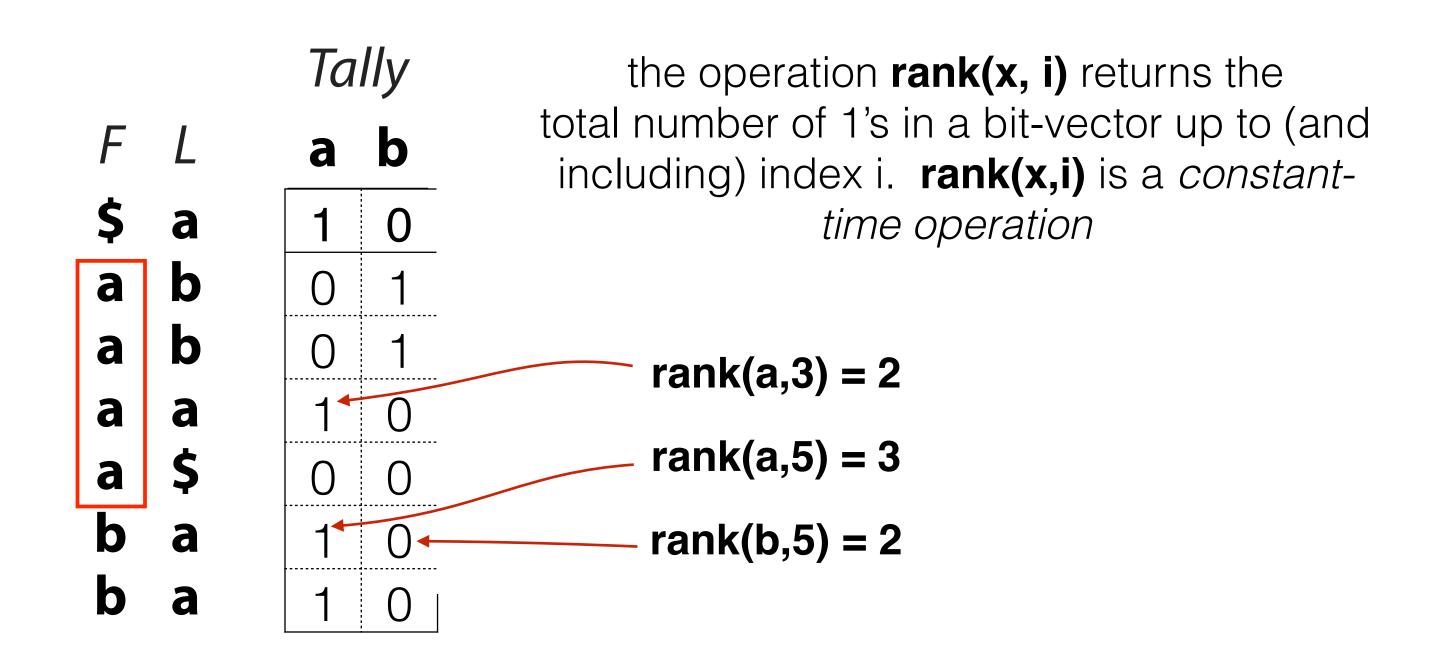
Another idea: pre-calculate # \mathbf{a} s, \mathbf{b} s in L up to *some* rows, e.g. every 5th row. Call pre-calculated rows *checkpoints*.



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*.



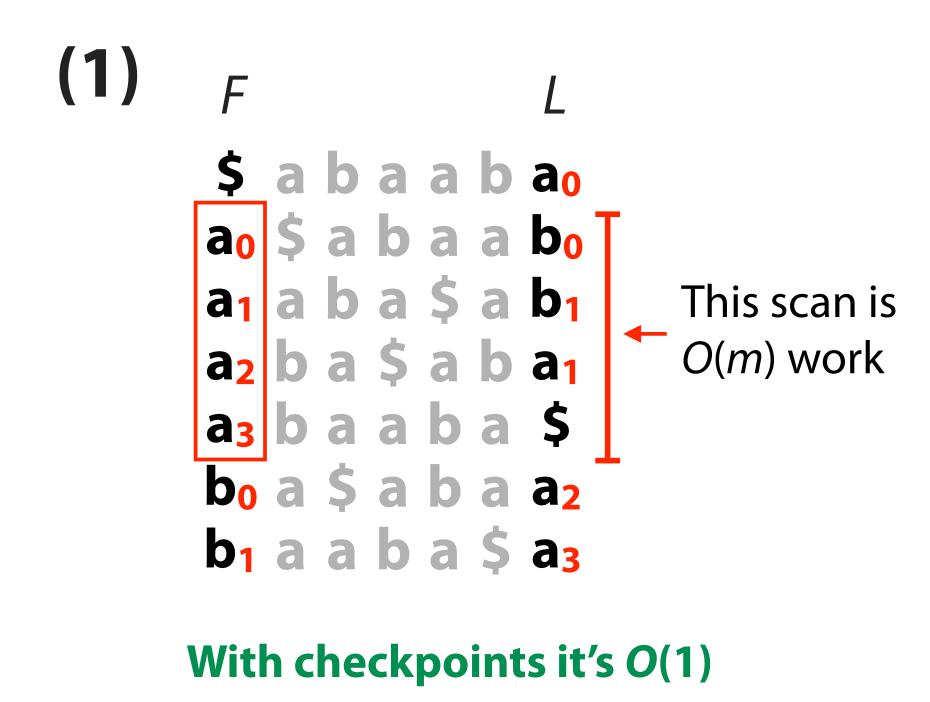
This can also be accomplished using **bit-vector rank** operations. We store one bit-vector for each character of Σ , placing a 1 where this character occurs and a 0 everywhere else:



To resolve the rank for a given character \mathbf{c} at a given index \mathbf{i} , we simply issue a $\mathbf{rank}(\mathbf{c},\mathbf{i})$ query. This is a practically-fast constant-time operation, but we need to keep around Σ bit-vectors, each of o(m) bits.

FM Index: a few problems

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



(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
```

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

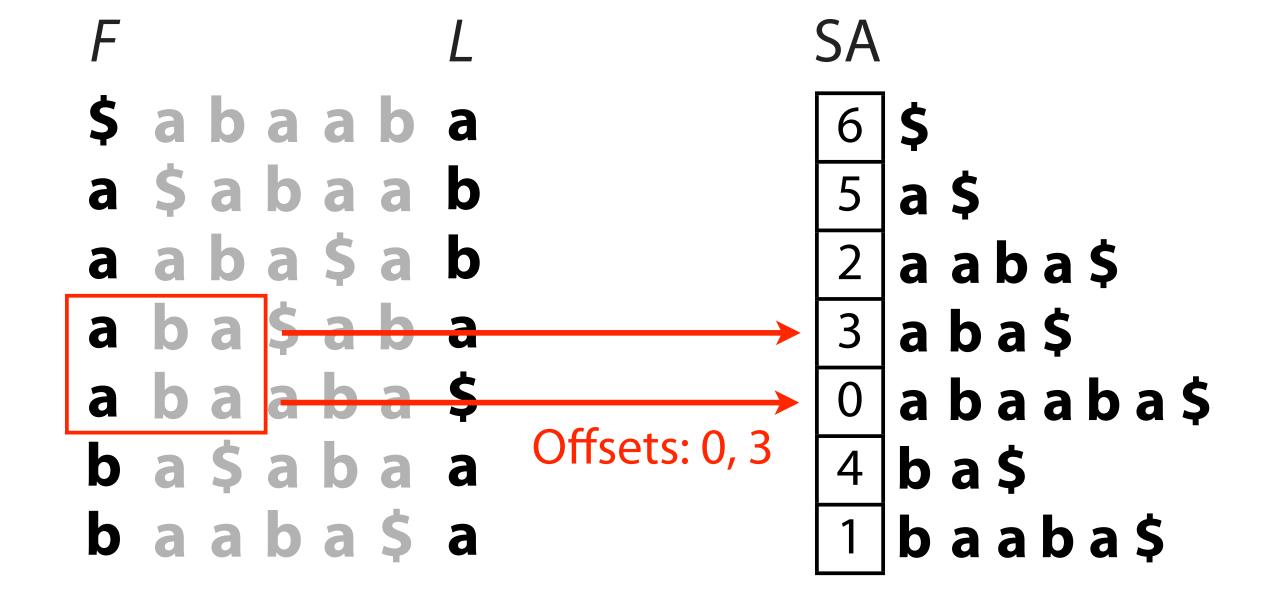
FM Index: a few problems

Not yet solved:

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

\$ ab a ab a₀
a₀ \$ ab a a b₀
a₁ ab a \$ a b₁
a₂ ba \$ ab a
a₃ ba ab a
b₀
a₁ ab a \$ ab a
b₁
a₃ ba ab a
a₂
b₁ aab a\$ a₃

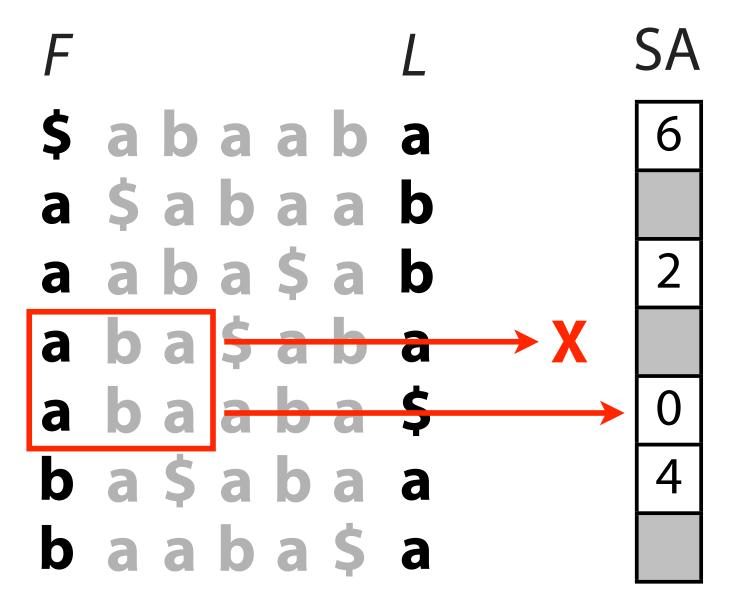
If suffix array were part of index, we could simply look up the offsets



But SA requires *m* integers

FM Index: resolving offsets

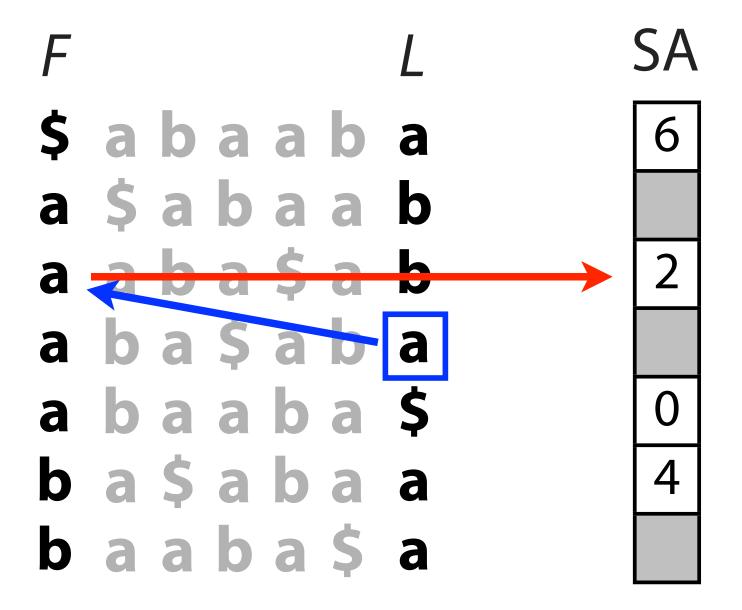
Idea: store some, but not all, entries of the suffix array



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 begining 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 a<sub>0</sub>
a<sub>0</sub> $ a b a a b a<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>
```

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 / $32 = \sim 400 \text{ MB}$

Checkpoints: 3 billion * 4 bytes/char * 4 char / 128 = ~400MB

Total ∼ 1.5 GB

Computing BWT in O(n) time

- Easy O(n² log n)-time algorithm to compute the BWT (create and sort the BWT matrix explicitly).
- Several direct O(n)-time algorithms for BWT.
 These are space efficient. (Bowtie e.g. uses [1])
- Also can use suffix arrays or trees:
 - Compute the suffix array, use correspondence between suffix array and BWT to output the BWT.
 - O(n)-time and O(n)-space, but the constants are large.

[1] Kärkkäinen, Juha. "Fast BWT in small space by blockwise suffix sorting." *Theoretical Computer Science* 387.3 (2007): 249-257.