Exact String Matching and searching for SNPs (2) CMSC423

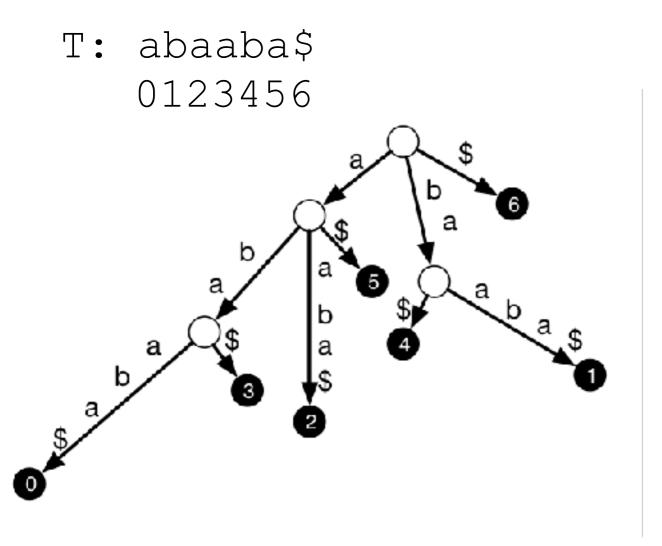
The problem

- Given:
 - 100's of millions of short reads: 100-200bp reads
 - A long reference genome (~3Bbp for human)
- Do:
 - Find high scoring scoring (fitting) alignments for each read
- What we know:
 - -Dynamic programming solution for fitting alignment:
 - 1e8 * 1e9 * 1e2 operations, 1e9 * 1e2 memory

Strategies

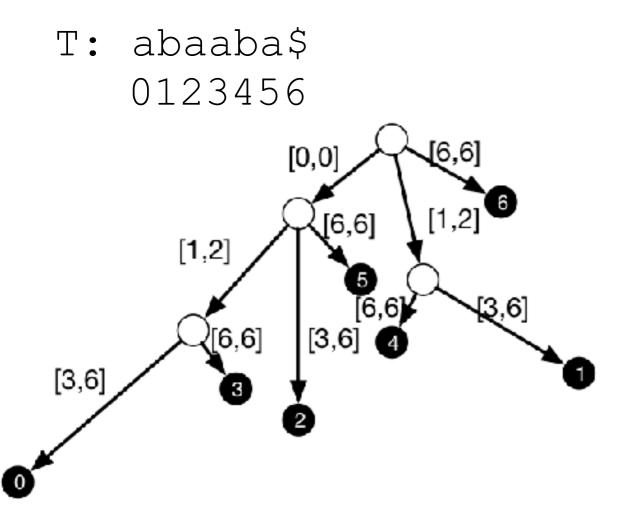
- What if we only allow a small number of substitutions?
 - Let's first try to find *exact* matches and work from those (the d+1 trick in the midterm)
- We are aligning to the same reference 100's of millions of time
 - —Is there preprocessing we can do to amortize time?
- Genomes are repetitive
 - -Can we search for matches in the genome in a smart way?
 - –Can we compress the genome, and search over the compressed representation?

Suffix Tree



- Collapse non-branching nodes
 - #nodes O(ITI)
- Memory requirement is *not* O(ITI)
 - In the worst case, space required for edge labels is O(ITI)

Suffix Tree



- Collapse non-branching nodes
 - #nodes O(ITI)
- Label edges with substring [start,end]
 O(1) per edge
- Memory now O(ITI)
- Construction algorithm O(ITI) (see Gusfield)

Recap

Structure	Processing Time	Memory	Search
Suffix Trie	O(T)	O(T ²)	O(P)
Suffix Tree	O(T)	O(T)*	O(P)
Suffix Array	O(T)	O(T) (but much smaller than Suffix Tree)	O(P log ₂ T)

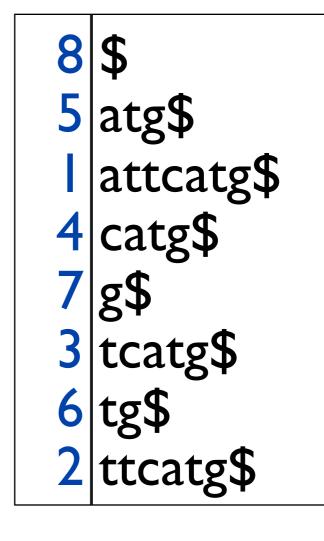
*In best implementations about 20 bytes per character (as opposed to 4 bytes for suffix array)

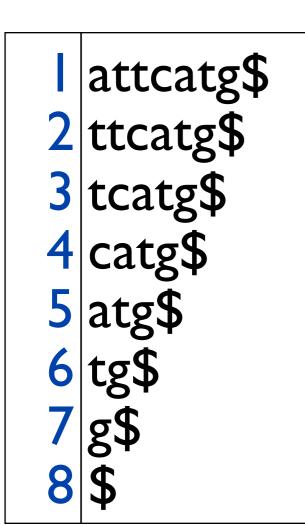
Suffix Arrays

- Even though Suffix Trees are O(n) space, the constant hidden by the big-Oh notation is somewhat "big": ≈ 20 bytes / character in good implementations.
- If you have a 10Gb genome, 20 bytes / character = 200Gb to store your suffix tree. "Linear" but large.
- Suffix arrays are a more efficient way to store the suffixes that can do most of what suffix trees can do, but just a bit slower.
- Slight space vs. time tradeoff.

Example Suffix Array

- Idea: lexicographically sort all the suffixes.
- Store the starting indices of the suffixes in an array.





suffix of s

sort the suffixes alphabetically

the indices just "come along for the ride"

index of suffix

s = attcatg

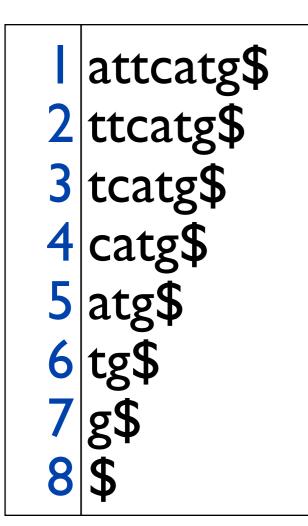
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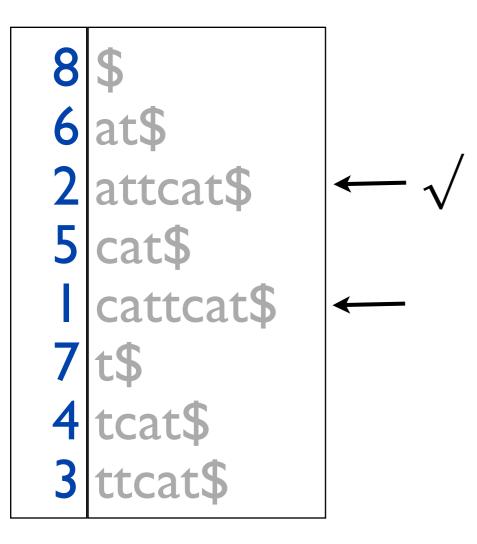
s = attcatg\$



index of suffix suffix of s

Search via Suffix Arrays

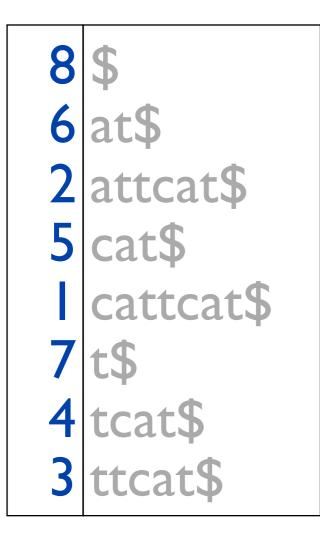
s = cattcat



- Does string "at" occur in s?
- Binary search to find "at".
- What about "tt"?

Counting via Suffix Arrays

s = cattcat\$



- How many times does "at" occur in the string?
- All the suffixes that start with "at" will be next to each other in the array.
- Find one suffix that starts with "at" (using binary search).
- Then count the neighboring sequences that start with at.

Constructing Suffix Arrays

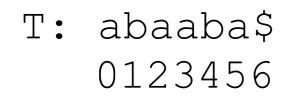
• Easy O(n² log n) algorithm:

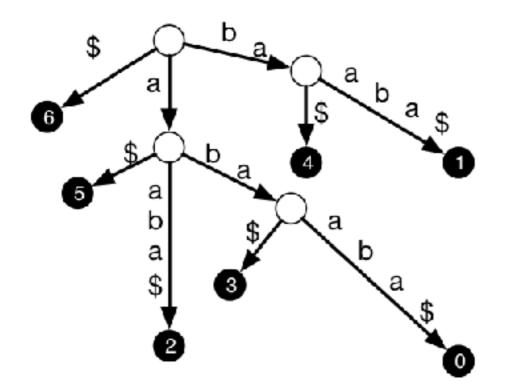
sort the n suffixes, which takes $O(n \log n)$ comparisons, where each comparison takes O(n).

• There are several direct O(n) algorithms for constructing suffix arrays that use very little space.

• An simple O(n) algorithm: build the suffix tree, and exploit the relationship between suffix trees and suffix arrays (next slide)

Relationship between Suffix Arrays and Suffix Trees





6	\$
5	a\$
2	aaba\$
3	aba\$
0	abaaba\$
4	ba\$
1	baaa\$

Build suffix trees with edge labels sorted lexicographically Order of leaves: 6,5,2,3,0,4,1

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*In best implementations about 20 bytes per character (as opposed to 4 bytes for suffix array)

Burrows-Wheeler Transform

Text transform that is useful for compression & search.

banana banana\$ \$banana a\$banan anana\$b nana\$ba ana\$ban sort ana\$ban anana\$b na\$bana banana\$ a\$banan nana\$ba \$banana na\$bana

BWT(banana) = annb\$aa

Tends to put runs of the same character together.

Makes compression work well.

"bzip" is based on this.

Another Example

appellee\$ appellee\$ \$appellee appellee\$ ppellee\$a pellee\$ap e\$appelle ellee\$app ee\$appell sort llee\$appe ellee\$app lee\$appel lee\$appel ee\$appell llee\$appe pellee\$ap e\$appelle ppellee\$a \$appellee

BWT(appellee\$) = e\$elplepa

Doesn't always improve the compressibility...

1) Compute Burrows-Wheeler Transform of

abbaaba\$

Recovering the string * first 3 columns sort , first column A first 2 columns BWT BWT \$a \$ e\$a \$ap ė \$appellee \$ ap \$ap a app appellee\$ e\$ e\$a e\$appelle ee\$ e e ee\$appell Sort prepend sort ee ee\$ ee e these $3 \rightarrow$ ellee\$app BWT these 2 \rightarrow el lee\$appel columns column columns ell e Ρ e Ρ llee\$appe le le lee pellee\$ap Π ppellee\$a lle e II e pe pel **p**pe Ρ Ρ PP a p ppe a pp

Inverse BWT

```
def inverseBWT(s):
    B = [s<sub>1</sub>, s<sub>2</sub>, s<sub>3</sub>, ..., s<sub>n</sub>]
    for i = 1..n:
        sort B
        prepend s<sub>i</sub> to B[i]
    return row of B that ends with $
```

Another BWT Example

dogwood\$ ogwood\$d gwood\$do wood\$dog <u>sort</u> ood\$dogw od\$dogwo d\$dogwoo \$dogwood

\$dogwood d\$dogwoo dogwood\$ last column gwood\$do od\$dogwo ogwood\$d ood\$dogw wood\$dog

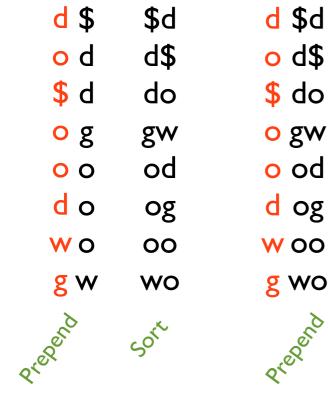
BWT(dogwood\$) = do\$oodwg

do\$oodwg Another BWT Example

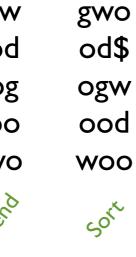
\$do

d\$d

dog











\$dog

d\$do

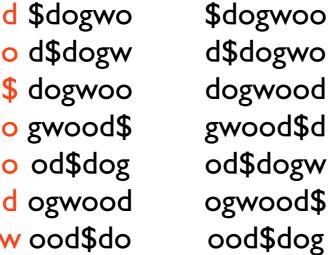
d\$do





wood\$do

Sort





Prepend

Sort

d \$dogw o d\$dog \$dogwo

- o gwood o od\$do
- d ogwoo
- wood\$d
- gwood\$

Prepend

od\$do d\$dog \$dogw dogwo gwood ogwoo ood\$d od\$do dogwo ogwoo ood\$d wood\$ wood\$ gwood

\$dogw

Solt

d \$dog



- d \$dogwoo
- d\$dogwo 0
- dogwood
- o gwood\$d
- o od\$dogw
- d ogwood\$
- w ood\$dog
- g wood\$do

Prepend

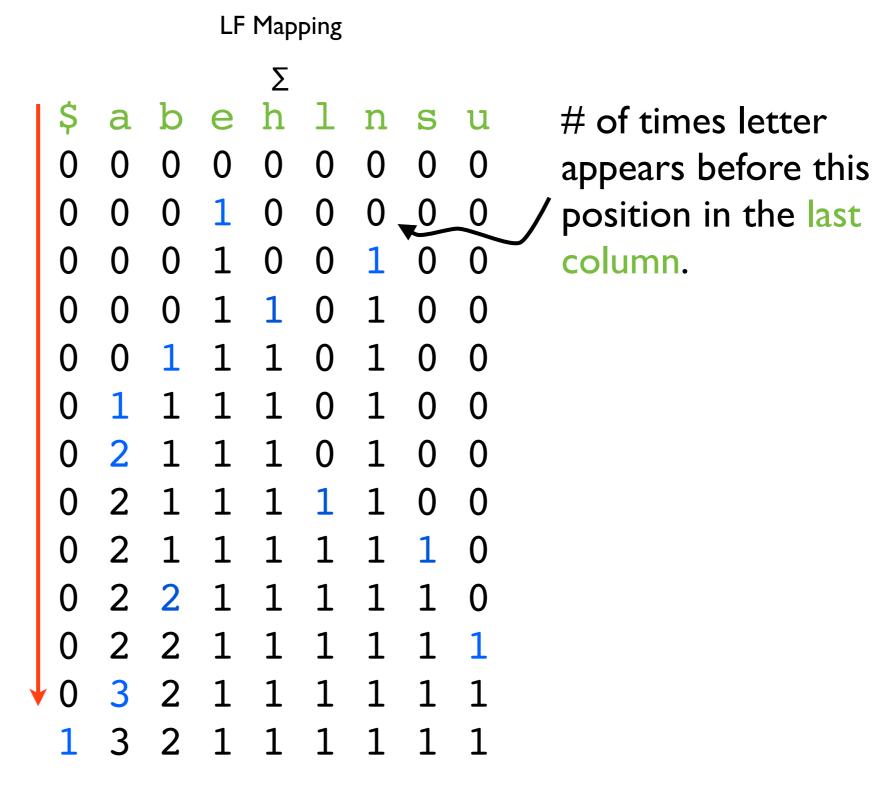
\$dogwood d\$dogwoo dogwood\$ gwood\$do od\$dogwo ogwood\$d ood\$dogw wood\$dog

Sort

Searching with BWT: LF Mapping

BWT(unabashable)

\$unabashable abashable\$un able\$unabash ashable\$unab bashable\$una ble\$unabasha e\$unabashabl hable\$unabas e\$unabashab nabashable\$u shable\$unaba unabashable\$



LF Property: The ith occurrence of a letter X in the last column corresponds to the ith occurrence of X in the first column.

1) Compute Burrows-Wheeler Transform of abbaaba\$

2) Write down the LF mapping matrix for the BWT of question 1

BWT Search

 $D(\Lambda/T/uncheckele)$

BWTSearch(aba) Start from the **end** of the pattern

Step I: Find the range of "a"s in the first column

Step 2: Look at the same range in the last column.

Step 3:"b" is the next
pattern character. Set B =
the LF mapping entry for b
in the first row of the
range.
Set E = the LF mapping
entry for b in the last + I
row of the range.

Step 4: Find the range for "b" in the first row, and use B and E to find the right subrange within the "b" range.

BWT(unabashable)
\$unabashable
abashable\$un
<mark>a</mark> ble\$unabash
ashable\$unab
bashable\$una
b le\$unabasha
e\$unabashabl
hable\$unabas
e\$unabashab
nabashable\$u
shable\$unaba
unabashable\$
-

LF Mapping

				Σ				
\$	a	b	е	h	1	n	S	u
0	0	0	0	0	0	0	0	0
0	0	0	1	0	0	0	0	0
0	0	0	1	0	0	1	0	0
0	0	0	1	1	0	1	0	0
0	0	1	1	1	0	1	0	0
0	1	1	1	1	0	1	0	0
0	2	1	1	1	0	1	0	0
0	2	1	1	1	1	1	0	0
0	2	1	1	1	1	1	1	0
0	2	2	1	1	1	1	1	0
0	2	2	1	1	1	1	1	1
0	3	2	1	1	1	1	1	1
1	3	2	1	1	1	1	1	1

BWT Searching Example 2

а	\$ a	b n
\$bananna	00	00
→ <mark>a</mark> \$banann	0	00
ananna \$b	0	0
anna\$ban	0	11
bananna\$	0	12
na\$banan		12
<mark>n</mark> anna \$ba		3
nna\$bana		13
	3	3
a Spananna		b n
<pre>\$bananna</pre>	00	00
<pre>\$bananna a\$banann</pre>		00000
\$bananna a\$banann ananna\$b	00	00
\$bananna a\$banann ananna\$b anna\$ban	00	00000
\$bananna a\$banann ananna\$b	00101	000 00 01 11
\$bananna a\$banann ananna\$b anna\$ban bananna\$	00101	000000000000000000000000000000000000000
<pre>\$bananna a\$bananna\$b anana\$ban bananna\$ ha\$banana\$</pre>	00 01 01 01 01 01	00 00 01 11 12 12

n \$bananna a\$banann ananna\$b anna\$ban bananna\$ bananna\$	0 0 0 0 0 0 1 0 2	₹ a a b →
nanna\$ba nna\$bana (B,E) = a	3 2 3 3 3	→n →
<pre>\$bananna a\$bananna\$b anna\$ban bananna\$ban bananna\$ba na\$banan nanna\$ba na\$banaa (B,E) = 0, I</pre>	0 0 0 0 0 1 0 0 0 1 0 1 0 1 1 1 0 1 1 2 1 1 1 2 1 1 1 3 1 2 1 3 1 3 1 3	-

pattern = "bana"

 \rightarrow

b n	n <mark>\$</mark> abn
00	\$ bananna 0000
00	a\$banann 0100
0	ananna\$b 0 0
11	anna\$ban 0
12	bananna 0112
12	na\$banan 2
3	nanna\$ba 3
13	nna\$bana 2 3
13	

Ь	\$ a b n
\$bananna	0000
<mark>a</mark> \$banann	0100
ananna \$b	0 0
<mark>a</mark> nna\$ban	0
bananna\$	0112
na\$banan	2
nanna\$ba	3
nna\$bana	2 3
	3 3

BWT Searching Notes

- Don't have to store the LF mapping. A more complex algorithm (later slides) lets you compute it in O(1) time in *compressed* data on the fly with some extra storage.
- To find the range in the first column corresponding to a character:
 - Pre-compute array C[c] = # of occurrences in the string of characters lexicographically < c.
 - Then start of the "a" range, for example, is: C["a"] + 1.
- Running time: O(|pattern|)
 - Finding the range in the first column takes O(I) time using the C array.
 - Updating the range takes O(I) time using the LF mapping.

Pseudocode for CountingOccurrences in BWT w/o stored LF mapping

C[c] = index into first column function Count(S_{bwt}, P): where the "c"s begin. c = P[p], i = psp = C[c] + 1; ep = C[c+1]while $(sp \le ep)$ and $(i \ge 2)$ do c = P[i-1]sp = C[c] + Occ(c, sp-1) + 1ep = C[c] + Occ(c, ep)i = i - 1 $-\mathbf{Occ}(c, p) = \# \text{ of of } c \text{ in the}$ first p characters of BWT(S), if ep < sp then aka the LF mapping. return "not found" else return ep - sp + 1

Relationship Between s = appellee\$ 123456789 **BWT and Suffix Arrays**

9

\$appellee appellee\$ e\$appelle ee\$appell ellee\$app lee\$appel llee\$appe pellee\$ap ppellee\$a

> BWT matrix

\$ appellee\$ e\$ ee\$ ellee\$ lee\$ llee\$ pellee\$ ppellee\$

The suffixes

are obtained

by deleting

everything

after the \$

These are still in sorted order because "\$" comes before everything else

8 7 - subtract $1 \rightarrow s[4-1] = p$ 4 6 5 3

Suffix array (start position for the suffixes) Suffix position - I =the position of the last character of the BWT matrix

s[9-1] = e

s[|-|] = \$

s[8-1] = e

s[7-1] = 1

s[6-1] = 1

s[5-1] = e

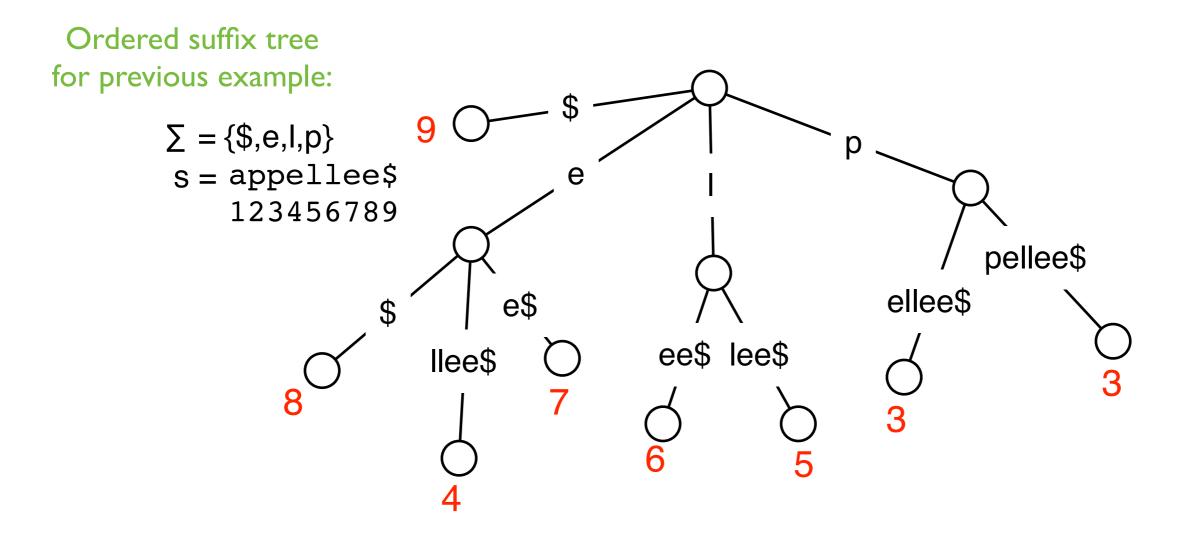
s[3-1] = p

s[2-1] = a

(\$ is a special case)

Relationship Between BWT and Suffix Trees

- Remember: Suffix Array = suffix numbers obtained by traversing the leaf nodes of the (ordered) Suffix Tree from left to right.
- Suffix Tree \Rightarrow Suffix Array \Rightarrow BWT.



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.
- 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.

Recap

BWT useful for searching and compression.

BWT is *invertible*: given the BWT of a string, the string can be reconstructed!

BWT is computable in O(n) time.

Close relationships between Suffix Trees, Suffix Arrays, and BWT:

- Suffix array = order of the suffix numbers of the suffix tree, traversed left to right
- BWT = letters at positions given by the suffix array entries I

Even after compression, can search string quickly.

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Structure	Processing Time	Memory	Search
Suffix Trie	O(T)	O(T ²)	O(P)
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Suffix Array	O(T)	O(T) (but much smaller than Suffix Tree)	O(P log ₂ T)
BWT	O(T)	O(T)**	O(P)

*In best implementations about 20 bytes per character (as opposed to 4 bytes for suffix array) **Compressed! For human genome ~2GB

Move-To-Front Coding

To encode a letter, use its index in the current list, and then move it to the front of the list.

	Σ	do\$oodwg
	\$dgow	1
List with all	d\$gow	13
letters from the allowed alphabet	od\$gw	132
	\$odgw	1322
	o\$dgw	13220
	o\$dgw	132202
	do\$gw	1322024
	wdo\$g	$13220244 = MTF(do \circ)$

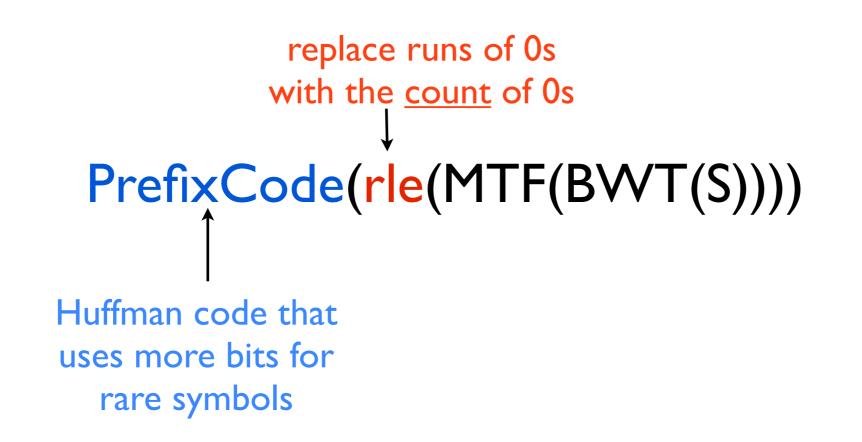
Benefits:

- Runs of the same letter will lead to runs of 0s.
- Common letters get small numbers, while rare letters get big numbers.

Compressing BWT Strings

Lots of possible compression schemes will benefit from preprocessing with BWT (since it tends to group runs of the same letters together).

One good scheme proposed by Ferragina & Manzini:

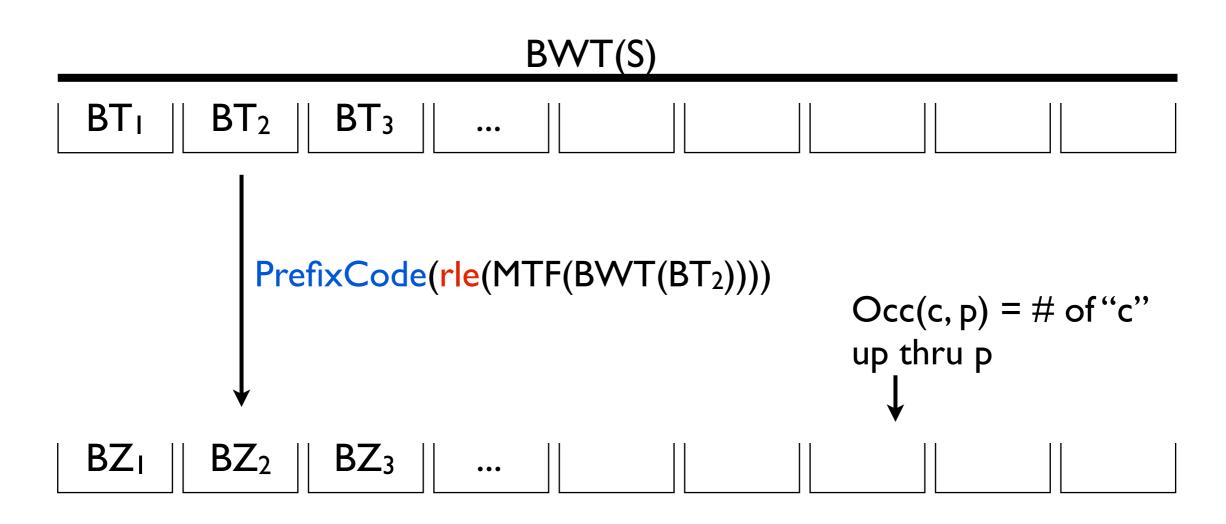


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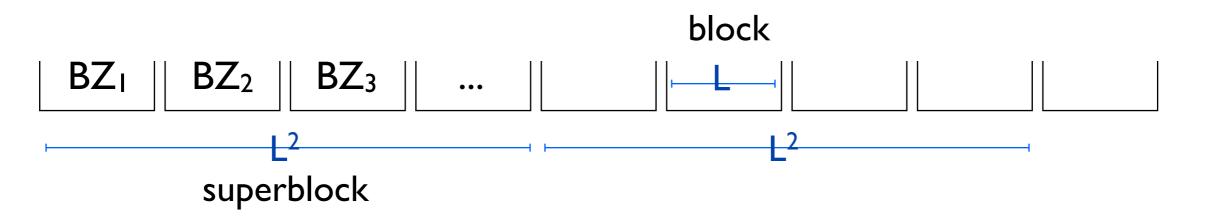
Computing Occ in Compressed String

Break BWT(S) into blocks of length L (we will decide on a value for L later):



Assumes every run of 0s is contained in a block [just for ease of explanation]. We will store some extra info for each block (and some groups of blocks) to compute Occ(c, p) quickly.

block: store $|\Sigma|$ -long array giving # of occurrences of each character up thru and including this block since the end of the last super block.

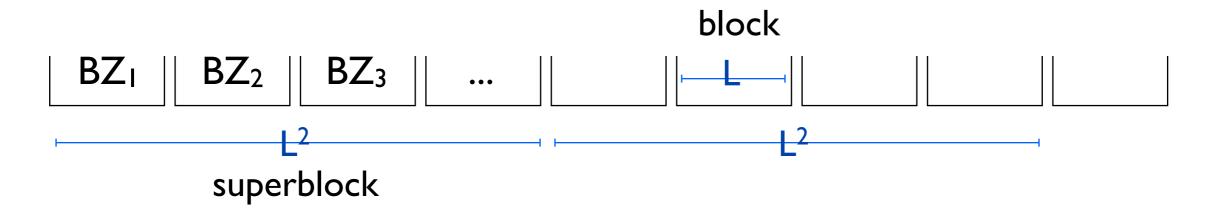


superblock: store $|\Sigma|$ -long array giving # of occurrences of each character up thru *and including* this superblock

u = compressed lengthChoose L = O(log u)

u/L blocks, each array is $\sum \log L \log x$ and $\sum \frac{u}{\log L} \log L = \frac{u}{\log u} \log \log u$ total space.

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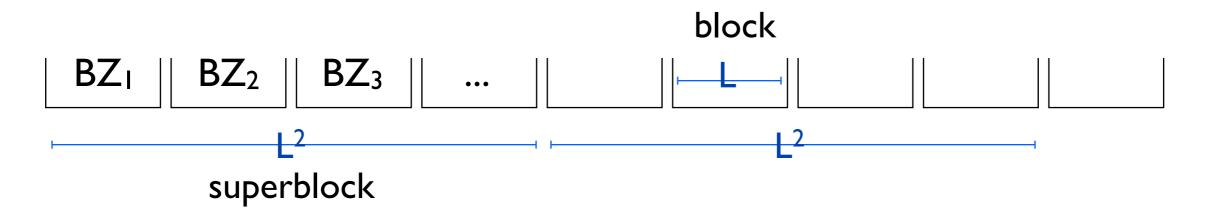


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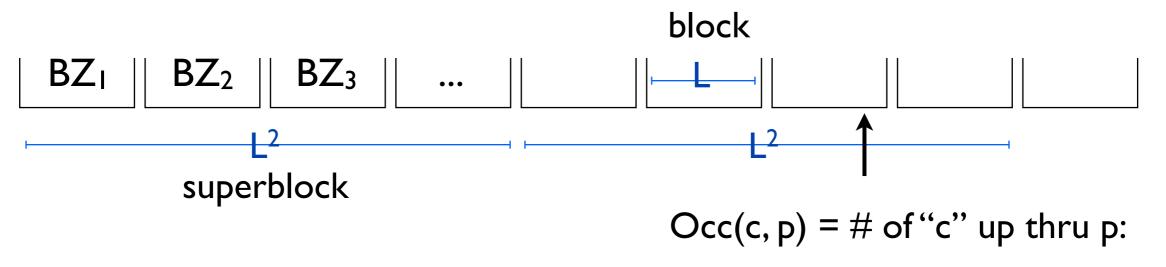
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superblock: store $|\Sigma|$ -long array giving # of occurrences of each character up thru *and including* this superblock

u/L² superblocks, each array is $|\sum \log u|$ long $\Rightarrow \frac{u}{(\log u)^2} \log u = \frac{u}{\log u}$ total space.

u = compressed lengthChoose L = O(log u)



sum value at last superblock, value at end of previous block, but then need to handle *this block*.

Store an array: $M[c, k, BZ_i, MTF_i] = #$ of occurrences of c through the kth letter of a block of <u>type</u> (BZ_i, MTF_i).

Size: $O(|\Sigma|L2^{L}|\Sigma|) = O(L2^{L'}) = O(u^{c}\log u)$ for c < 1 (since the string is compressed)