FM Index

Michael Schatz

Feb 18, 2013

Lecture 7: Applied Comparative Genomics



Assignment 3: Due Monday Feb 25

Assignment 3: Coverage, Genome Assembly, and the BWT

Assignment Date: Monday, Feb. 18, 2019 Due Date: Monday, Feb. 25, 2019 @ 11:59pm

Question 1. Coverage simulator [10 pts]

- Q1a. How many 100bp reads are needed to sequence a 1Mbp genome to 5x coverage?
- Q1b. In the language of your choice, simulate sequencing 5x coverage of a 1Mbp genome and plot the histogram of coverage. Note you do not need to actually output the
 sequences of the reads, you can just randomly sample positions in the genome and record the coverage. You do not need to consider the strand of each read. The start
 position of each read should have a uniform random probability at each possible starting position (1 through 999,900). You can record the coverage in an array of 1M
 positions. Overlay the histogram with a Poisson distribution with lambda=5
- Q1c. Using the histogram from 1b, how much of the genome has not been sequenced (has 0x coverage). How well does this match Poisson expectations?
- Q1d. Now repeat the analysis with 15x coverage: 1. simulate the appropriate number of reads, 2. make a histogram, 3. overlay a Poisson distribution with lambda=15, 4.
 compute the number of bases with 0x coverage, and 5. evaluated how well it matches the Poisson expectation.

Question 2. de Bruijn Graph construction [10 pts]

 Q1a. Draw (by hand or by code) the de Bruijn graph for the following reads using k=3 (assume all reads are from the forward strand, no sequencing errors, complete coverage of the genome)

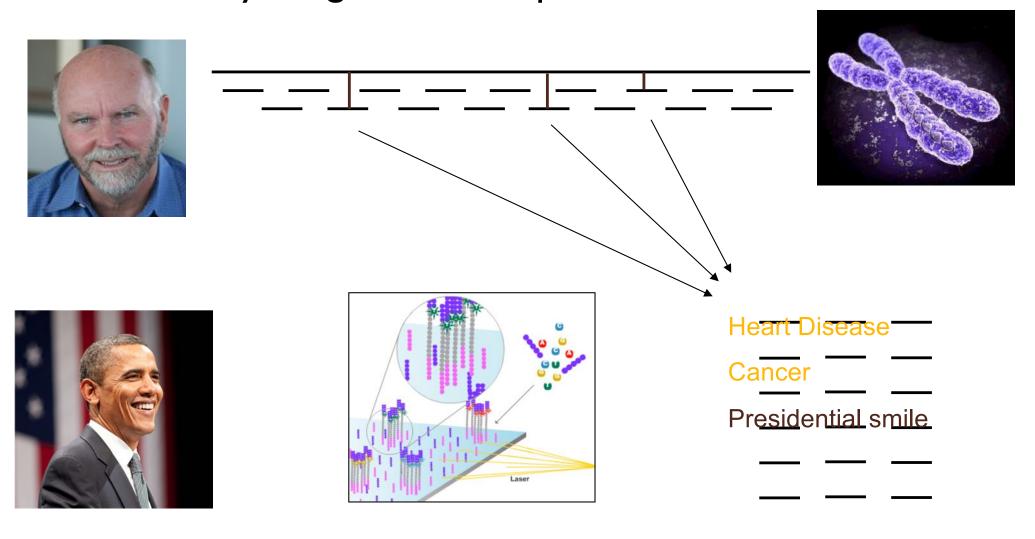
ATTC ATTG CATT CTTA GATT TATT TCAT TCTT



Part I: Suffix Arrays

Personal Genomics

How does your genome compare to the reference?



Brute Force Analysis



- Brute Force:
 - At every possible offset in the genome:
 - Do all of the characters of the query match?
- Analysis
 - Simple, easy to understand

Genome length = n	[3B]
– Query length = m	[7]
Comparisons: (n-m+1) * m	[21B]

Overall runtime: O(nm)

[How long would it take if we double the genome size, read length?] [How long would it take if we double both?]

Brute Force Reflections

Why check every position?

- GATTACA can't possibly start at position 15

[WHY?]

I	2	3	4	5	6	7	8	9	10	П	12	13	14	15	•••
Т	G	Α	Т	Т	Α	С	Α	G	Α	Т	Т	Α	С	С	• • •
								G	Α	Т	Т	Α	С	Α	

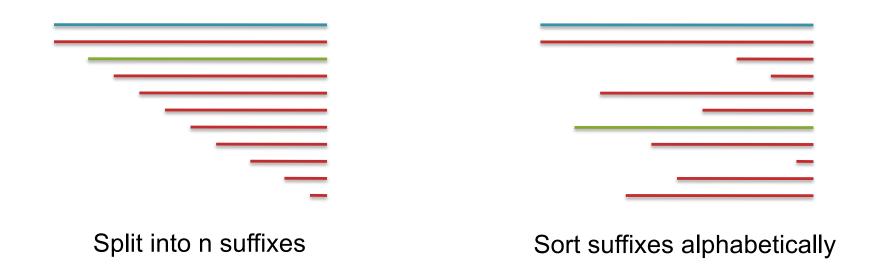
- Improve runtime to O(n + m)

[3B + 7]

- If we double both, it just takes twice as long
- Knuth-Morris-Pratt, 1977
- Boyer-Moyer, 1977, 1991
- For one-off scans, this is the best we can do (optimal performance)
 - We have to read every character of the genome, and every character of the query
 - For short queries, runtime is dominated by the length of the genome

Suffix Arrays: Searching the Phone Book

- What if we need to check many queries?
 - We don't need to check every page of the phone book to find 'Schatz'
 - Sorting alphabetically lets us immediately skip 96% (25/26) of the book without any loss in accuracy
- Sorting the genome: Suffix Array (Manber & Myers, 1991)
 - Sort every suffix of the genome



[Challenge Question: How else could we split the genome?]

- Strategy 2: Binary search
 - Compare to the middle, refine as higher or lower
- Searching for GATTACA
 - Lo = I; Hi = 15;



#	Sequence	Pos
_	ACAGATTACC	6
2	ACC	13
3	AGATTACC	8
4	ATTACAGATTACC	3
5	ATTACC	10
6	C	15
7	CAGATTACC	7
8	CC	14
9	GATTACAGATTACC	2
10	GATTACC	9
П	TACAGATTACC	5
12	TACC	12
13	TGATTACAGATTACC	I
14	TTACAGATTACC	4
15	TTACC	Ш



- Strategy 2: Binary search
 - Compare to the middle, refine as higher or lower
- Searching for GATTACA
 - Lo = I; Hi = 15; Mid = (1+15)/2 = 8
 - Middle = Suffix[8] = CC



#	Sequence	Pos
I	ACAGATTACC	6
2	ACC	13
3	AGATTACC	8
4	ATTACAGATTACC	3
5	ATTACC	10
6	C	15
7	CAGATTACC	7
8	CC	14
9	GATTACAGATTACC	2
10	GATTACC	9
П	TACAGATTACC	5
12	TACC	12
13	TGATTACAGATTACC	I
14	TTACAGATTACC	4
15	TTACC	Ш



- Strategy 2: Binary search
 - Compare to the middle, refine as higher or lower
- Searching for GATTACA
 - Lo = I; Hi = 15; Mid = (1+15)/2 = 8
 - Middle = Suffix[8] = CC=> Higher: Lo = Mid + I



#	Sequence	Pos
I	ACAGATTACC	6
2	ACC	13
3	AGATTACC	8
4	ATTACAGATTACC	3
5	ATTACC	10
6	C	15
7	CAGATTACC	7
8	CC	14
9	GATTACAGATTACC	2
10	GATTACC	9
П	TACAGATTACC	5
12	TACC	12
13	TGATTACAGATTACC	I
14	TTACAGATTACC	4
15	TTACC	Ш



- Strategy 2: Binary search
 - Compare to the middle, refine as higher or lower
- Searching for GATTACA
 - Lo = I; Hi = 15; Mid = (1+15)/2 = 8
 - Middle = Suffix[8] = CC=> Higher: Lo = Mid + I
 - Lo = 9; Hi = 15;

3	AGATTACC	8
4	ATTACAGATTACC	3
5	ATTACC	10
6	C	15
7	CAGATTACC	7
8	CC	14
9	GATTACAGATTACC	2
10	GATTACC	9
П	TACAGATTACC	5
12	TACC	12
	1400	12

TGATTACAGATTACC...

TTACAGATTACC...

15 TTACC...

Pos

13

4

П

Sequence

ACC...

ACAGATTACC...





- Strategy 2: Binary search
 - Compare to the middle, refine as higher or lower
- Searching for GATTACA
 - Lo = I; Hi = 15; Mid = (1+15)/2 = 8
 - Middle = Suffix[8] = CC=> Higher: Lo = Mid + I
 - Lo = 9; Hi = 15; Mid = (9+15)/2 = 12
 - Middle = Suffix[12] = TACC

#	Sequence	Pos
- 1	ACAGATTACC	6
2	ACC	13
3	AGATTACC	8
4	ATTACAGATTACC	3
5	ATTACC	10
6	C	15
7	CAGATTACC	7
8	CC	14
9	GATTACAGATTACC	2
10	GATTACC	9
П	TACAGATTACC	5
12	TACC	12
13	TGATTACAGATTACC	I
14	TTACAGATTACC	4
15	TTACC	Ш





- Strategy 2: Binary search
 - Compare to the middle, refine as higher or lower
- Searching for GATTACA
 - Lo = I; Hi = 15; Mid = (1+15)/2 = 8
 - Middle = Suffix[8] = CC=> Higher: Lo = Mid + I
 - Lo = 9; Hi = 15; Mid = (9+15)/2 = 12
 - Middle = Suffix[12] = TACC=> Lower: Hi = Mid I
 - Lo = 9; Hi = 11;





#	Sequence	Pos
- 1	ACAGATTACC	6
2	ACC	13
3	AGATTACC	8
4	ATTACAGATTACC	3
5	ATTACC	10
6	C	15
7	CAGATTACC	7
8	CC	14
9	GATTACAGATTACC	2
10	GATTACC	9
П	TACAGATTACC	5
12	TACC	12
13	TGATTACAGATTACC	I
14	TTACAGATTACC	4
15	TTACC	П

- Strategy 2: Binary search
 - Compare to the middle, refine as higher or lower
- Searching for GATTACA
 - Lo = I; Hi = 15; Mid = (1+15)/2 = 8
 - Middle = Suffix[8] = CC=> Higher: Lo = Mid + I
 - Lo = 9; Hi = 15; Mid = (9+15)/2 = 12
 - Middle = Suffix[12] = TACC=> Lower: Hi = Mid I
 - Lo = 9; Hi = 11; Mid = (9+11)/2 = 10
 - Middle = Suffix[10] = GATTACC

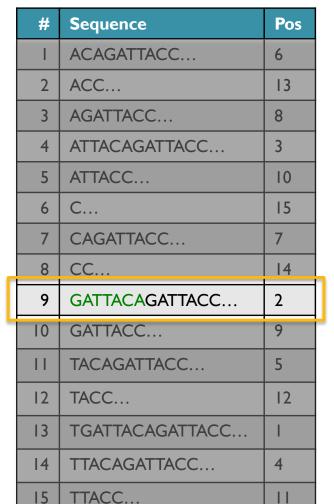
#	Sequence	Pos
I	ACAGATTACC	6
2	ACC	13
3	AGATTACC	8
4	ATTACAGATTACC	3
5	ATTACC	10
6	C	15
7	CAGATTACC	7
8	CC	14
9	GATTACAGATTACC	2
10	GATTACC	9
П	TACAGATTACC	5
12	TACC	12
13	TGATTACAGATTACC	I
14	TTACAGATTACC	4
15	TTACC	Ш

- Strategy 2: Binary search
 - Compare to the middle, refine as higher or lower
- Searching for GATTACA
 - Lo = I; Hi = 15; Mid = (1+15)/2 = 8
 - Middle = Suffix[8] = CC=> Higher: Lo = Mid + I
 - Lo = 9; Hi = 15; Mid = (9+15)/2 = 12
 - Middle = Suffix[12] = TACC=> Lower: Hi = Mid I
 - Lo = 9; Hi = 11; Mid = (9+11)/2 = 10
 - Middle = Suffix[10] = GATTACC=> Lower: Hi = Mid I
 - Lo = 9; Hi = 9;



#	Sequence	Pos
_	ACAGATTACC	6
2	ACC	13
3	AGATTACC	8
4	ATTACAGATTACC	3
5	ATTACC	10
6	C	15
7	CAGATTACC	7
8	CC	14
9	GATTACAGATTACC	2
10	GATTACC	9
П	TACAGATTACC	5
12	TACC	12
13	TGATTACAGATTACC	I
14	TTACAGATTACC	4
15	TTACC	П

- Strategy 2: Binary search
 - Compare to the middle, refine as higher or lower
- Searching for GATTACA
 - Lo = I; Hi = 15; Mid = (1+15)/2 = 8
 - Middle = Suffix[8] = CC=> Higher: Lo = Mid + I
 - Lo = 9; Hi = 15; Mid = (9+15)/2 = 12
 - Middle = Suffix[12] = TACC=> Lower: Hi = Mid I
 - Lo = 9; Hi = 11; Mid = (9+11)/2 = 10
 - Middle = Suffix[10] = GATTACC=> Lower: Hi = Mid I
 - Lo = 9; Hi = 9; Mid = (9+9)/2 = 9
 - Middle = Suffix[9] = GATTACA...=> Match at position 2!





Binary Search Analysis

Binary Search

```
Initialize search range to entire list

mid = (hi+lo)/2; middle = suffix[mid]

if query matches middle: done

else if query < middle: pick low range

else if query > middle: pick hi range

Repeat until done or empty range
```

[WHEN?]

- Analysis
 - More complicated method
 - How many times do we repeat?
 - How many times can it cut the range in half?
 - Find smallest x such that: $n/(2^x) \le 1$; $x = \lg_2(n)$

[32]

- Total Runtime: O(m lg n)
 - More complicated, but much faster!
 - Looking up a query loops 32 times instead of 3B

[How long does it take to search 6B or 24B nucleotides?]



Binary Search Analysis

Binary Search

```
Initialize search range to entire list

mid = (hi+lo)/2; middle = suffix[mid]

if query matches middle: done

else if query < middle: pick low range

else if query > middle: pick hi range

Repeat until done or empty range
```

[WHEN?]

[32]

- Analysis
 - More complicated method
 - How many times do we repeat?
 - How many times can it cut the range in half?
 - Find smallest x such that: $n/(2^x) \le 1$; $x = \lg_2(n)$
- Total Runtime: O(m lg n)
 - More complicated, but much faster!
 - Looking up a query loops 32 times instead of 3B



Can be reduced to O(m + lg n) using an auxiliary data structure called the LCP array

Suffix Array Construction

How can we store the suffix array?
 [How many characters are in all suffixes combined?]

$$S = 1 + 2 + 3 + \dots + n = \sum_{i=1}^{n} i = \frac{n(n+1)}{2} = O(n^2)$$

- Hopeless to explicitly store 4.5 billion billion characters
- Instead use implicit representation
 - Keep I copy of the genome, and a list of sorted offsets
 - Storing 3 billion offsets fits on a server (12GB)
- Searching the array is very fast, but it takes time to construct
 - This time will be amortized over many, many searches
 - Run it once "overnight" and save it away for all future queries



13

8

10

15

7

2

9

5

12

4

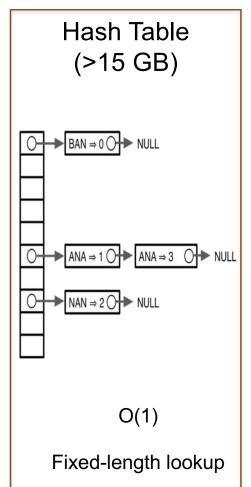
П

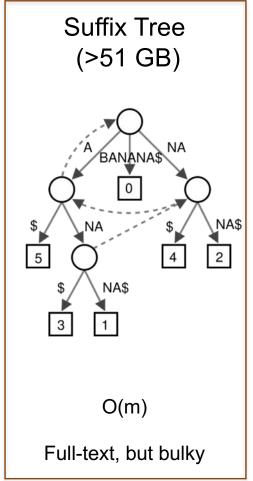
Exact Matching Review & Overview

Where is GATTACA in the human genome?

Brute Force (3 GB) BANANA BAN ANA NAN ANA O(m * n)Slow & Easy

Suffix Array (>15 GB) A\$ ANA\$ ANANA\$ BANANA\$ NA\$ NANA\$ $O(m + \lg n)$ Full-text index





*** These are general techniques applicable to any text search problem ***

Part 2: Burrows Wheeler Transform

Algorithmic challenge

How can we combine the speed of a suffix array O(m + lg(n)) (or even O(m)) with the size of a brute force analysis (n bytes)?

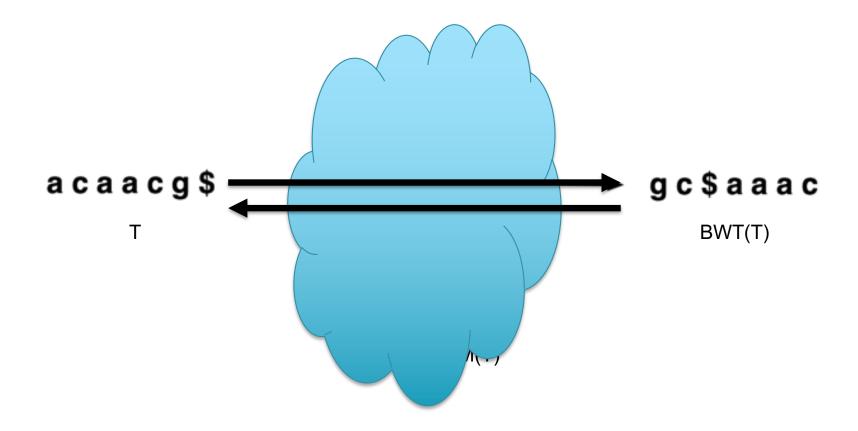
What would such an index look like?



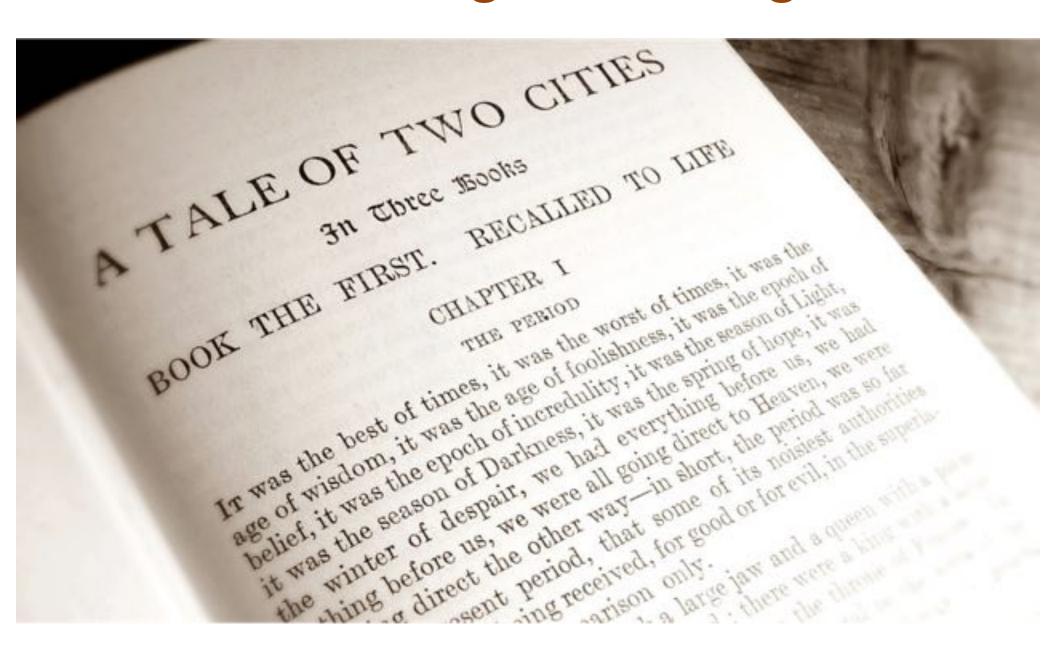
Bowtie: Ultrafast and memory efficient alignment of short DNA sequences to the human genome

Slides Courtesy of Ben Langmead

Reversible permutation of the characters in a text



A block sorting lossless data compression algorithm.



ref[614]:

It_was_the_best_of_times,_it_was_the_worst_of_times,_it_was_the_age_
of_wisdom,_it_was_the_age_of_foolishness,_it_was_the_epoch_of_belief
,_it_was_the_epoch_of_incredulity,_it_was_the_season_of_Light,_it_wa
s_the_season_of_Darkness,_it_was_the_spring_of_hope,_it_was_the_wint
er_of_despair,_we_had_everything_before_us,_we_had_nothing_before_us
,_we_were_all_going_direct_to_Heaven,_we_were_all_going_direct_the_o
ther_way_-_in_short,_the_period_was_so_far_like_the_present_period,_
that_some_of_its_noisiest_authorities_insisted_on_its_being_received
,_for_good_or_for_evil,_in_the_superlative_degree_of_comparison_only.\$

Run Length Encoding:

- Replace a "run" of a character X with a single X followed by the length of the run
- GAAAAAAATTACA => GA8T2ACA (reverse is also easy to implement)
- If your text contains numbers, then you will need to use a (slightly) more sophisticated encoding

ref[614]:

It_was_the_best_of_times,_it_was_the_worst_of_times,_it_was_the_age_
of_wisdom,_it_was_the_age_of_foolishness,_it_was_the_epoch_of_belief
,_it_was_the_epoch_of_incredulity,_it_was_the_season_of_Light,_it_wa
s_the_season_of_Darkness,_it_was_the_spring_of_hope,_it_was_the_wint
er_of_despair,_we_had_everything_before_us,_we_had_nothing_before_us
,_we_were_all_going_direct_to_Heaven,_we_were_all_going_direct_the_o
ther_way_-_in_short,_the_period_was_so_far_like_the_present_period,_
that_some_of_its_noisiest_authorities_insisted_on_its_being_received
,_for_good_or_for_evil,_in_the_superlative_degree_of_comparison_only.\$

rle(ref)[614]:

It_was_the_best_of_times,_it_was_the_worst_of_times,_it_was_the_age_
of_wisdom,_it_was_the_age_of_fo2lishnes2,_it_was_the_epoch_of_belief
,_it_was_the_epoch_of_incredulity,_it_was_the_season_of_Light,_it_wa
s_the_season_of_Darknes2,_it_was_the_spring_of_hope,_it_was_the_wint
er_of_despair,_we_had_everything_before_us,_we_had_nothing_before_us
,_we_were_al2_going_direct_to_Heaven,_we_were_al2_going_direct_the_o
ther_way_-_in_short,_the_period_was_so_far_like_the_present_period,_
that_some_of_its_noisiest_authorities_insisted_on_its_being_received
,_for_go2d_or_for_evil,_in_the_superlative_degre2_of_comparison_only.\$

ref[614]:

It_was_the_best_of_times,_it_was_the_worst_of_times,_it_was_the_age_
of_wisdom,_it_was_the_age_of_foolishness,_it_was_the_epoch_of_belief
,_it_was_the_epoch_of_incredulity,_it_was_the_season_of_Light,_it_wa
s_the_season_of_Darkness,_it_was_the_spring_of_hope,_it_was_the_wint
er_of_despair,_we_had_everything_before_us,_we_had_nothing_before_us
,_we_were_all_going_direct_to_Heaven,_we_were_all_going_direct_the_o
ther_way_-_in_short,_the_period_was_so_far_like_the_present_period,_
that_some_of_its_noisiest_authorities_insisted_on_its_being_received
,_for_good_or_for_evil,_in_the_superlative_degree_of_comparison_only.\$

bwt[614]:

bwt[614]:

<pre>.dlmssftysesdtrsns_y\$_yfofeeeetggsfefefggeedrofr,llreef-,fs,,,,,,</pre>
,,nfrsdnnhereghettedndeteegeenstee,ssssst,esssnssffteedtttttttttt,,
,,eeefehh_p_fpDwwwwwwwwwwwweehl_eweoo_neeeoaaeoosephhrrhvh
hwwegmghhhhhhhkrrwwhhssHrrrvtrribbdbcbvsthwwpppvmmirdnnibeoooooo
ooooooeennnnnaaiecctttttttttttttttttttts_tsgltsLlvtthhoor
e_wrraddwlorsr_lteirillre_ouaanooiioeooooiiihkiiiiiioiei
tsppioiggnodsc_sss_gfhf_fffhwh_nsmouee_sioooaeeeeoo_ii
cgppeeaoaeooeesseuutetaaaaaaaaaaiei_inaaie_eeerei_hrsssnacciiIi
iiiiiisnoyoui_a_iiids_aiiaeetlar

rle(bwt)[464]:

.dlms2ftysesdtrsns_y_2\$_yfofe4tg2sfefefg2e2drofr,12re2f-,fs,9nfrsdn2 hereghet2edndete2ge2nste2,s5t,es3ns2f2te2dt10r,4e3feh2_2p_2fpDw11e2h 1_ew_5eo2_ne3oa2eo2_4seph2r2hvh2w2egmgh7kr2w2h2s2Hr3vtr2ib2dbcbvs_2t hw2p3vm2irdn2ib_2eo12_4e2n6a2i_3ec2_2t18s_tsgltsLlvt2_3h2o2re_wr2ad2 wlors_9r_2lteiril2re_oua2no2i2oeo4i3hki6o_2ieitsp2ioi_12g2nodsc_s3_g fhf_f3hwh_nsmo_2ue2_sio3ae4o2_i2cgp2e2aoaeo2e2s2eu2teta11i_2ei_in_2a 2ie_e3rei_hrs3nac2i2Ii7sn_15oyoui_2a_i3ds_2ai2ae2_21tlar

bwt[614]:

rle(bwt)[464]:

.dlms2ftysesdtrsns_y_2\$_yfofe4tg2sfefefg2e2drofr,12re2f-,fs,9nfrsdn2 hereghet2edndete2ge2nste2,s5t,es3ns2f2te2dt10r,4e3feh2_2p_2fpDw11e2h 1_ew_5eo2_ne3oa2eo2_4seph2r2hvh2w2egmgh7kr2w2h2s2Hr3vtr2ib2dbcbvs_2t hw2p3vm2irdn2ib_2eo12_4e2n6a2i_3ec2_2t18s_tsgltsLlvt2_3h2o2re_wr2ad2 wlors_9r_2lteiril2re_oua2no2i2oeo4i3hki6o_2ieitsp2ioi_12g2nodsc_s3_g fhf_f3hwh_nsmo_2ue2_sio3ae4o2_i2cgp2e2aoaeo2e2s2eu2teta11i_2ei_in_2a 2ie_e3rei_hrs3nac2i2Ii7sn_15oyoui_2a_i3ds_2ai2ae2_21tlar

ref[614]:

It_was_the_best_of_times,_it_was_the_worst_of_times,_it_was_the_age_
of_wisdom,_it_was_the_age_of_foolishness,_it_was_the_epoch_of_belief
,_it_was_the_epoch_of_incredulity,_it_was_the_season_of_Light,_it_wa
s_the_season_of_Darkness,_it_was_the_spring_of_hope,_it_was_the_wint
er_of_despair,_we_had_everything_before_us,_we_had_nothing_before_us
,_we_were_all_going_direct_to_Heaven,_we_were_all_going_direct_the_o
ther_way_-_in_short,_the_period_was_so_far_like_the_present_period,_
that_some_of_its_noisiest_authorities_insisted_on_its_being_received
,_for_good_or_for_evil,_in_the_superlative_degree_of_comparison_only.\$

rle(bwt)[464]:

.dlms2ftysesdtrsns_y_2\$_yfofe4tg2sfefefg2e2drofr,12re2f-,fs,9nfrsdn2 hereghet2edndete2ge2nste2,s5t,es3ns2f2te2dt10r,4e3feh2_2p_2fpDw11e2h 1_ew_5eo2_ne3oa2eo2_4seph2r2hvh2w2egmgh7kr2w2h2s2Hr3vtr2ib2dbcbvs_2t hw2p3vm2irdn2ib_2eo12_4e2n6a2i_3ec2_2t18s_tsgltsLlvt2_3h2o2re_wr2ad2 wlors_9r_2lteiril2re_oua2no2i2oeo4i3hki6o_2ieitsp2ioi_12g2nodsc_s3_g fhf_f3hwh_nsmo_2ue2_sio3ae4o2_i2cgp2e2aoaeo2e2s2eu2teta11i_2ei_in_2a 2ie_e3rei_hrs3nac2i2Ii7sn_15oyoui_2a_i3ds_2ai2ae2_21tlar

ref[614]:

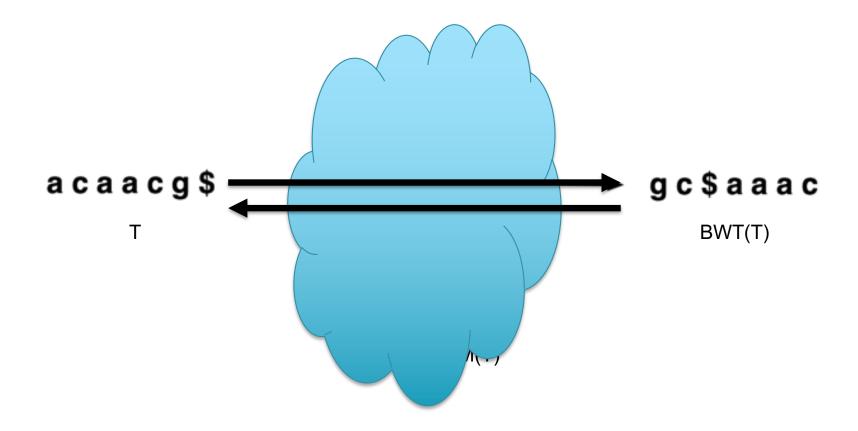
It_was_the_best_of_times,_it_was_the_worst_of_times,_it_was_the_age_
of_wisdom,_it_was_the_age_of_foolishness,_it_was_the_epoch_of_belief
,_it_was_the_epoch_of_incredulity,_it_was_the_season_of_Light,_it_wa
s_the_season_of_Darkness,_it_was_the_spring_of_hope,_it_was_the_wint
er_of_despair,_we_had_everything_before_us,_we_had_nothing_before_us
,_we_were_all_going_direct_to_Heaven,_we_were_all_going_direct_the_o
ther_way_-_in_short,_the_period_was_so_far_like_the_present_period,_
that_some_of_its_noisiest_authorities_insisted_on_its_being_received
,_for_good_or_for_evil,_in_the_superlative_degree_of_comparison_only.\$

rle(bwt)[464]:

.dlms2ftysesdtrsns_y_2\$_yfofe4tg2sfefefg2e2drofr,l2re2f-,fs,9nfrsdn2hereghet2edndete2ge2nste2,s5t,es3ns2f2te2dt10r,4e3feh2_2p_2fpDw11e2hl_ew_5eo2_ne3oa2eo2_4seph2r2hvh2w2egmgh7kr2w2h2s2Hr3vtr2ib2dbcbvs_2thw2p3vm2irdn2ib_2eo12_4e2n6a2i_3ec2_2t18s_tsgltsLlvt2_3h2o2re_wr2ad2wlors_9r_2lteiril2re_oua2no2i2oeo4i3hki6o_2ieitsp2ioi_12g2nodsc_s3_gfhf_f3hwh_nsmo_2ue2_sio3ae4o2_i2cgp2e2aoaeo2e2s2eu2teta11i_2ei_in_2a2ie_e3rei_Saved 614-464 = 150 bytes (24%) with zero loss of information!

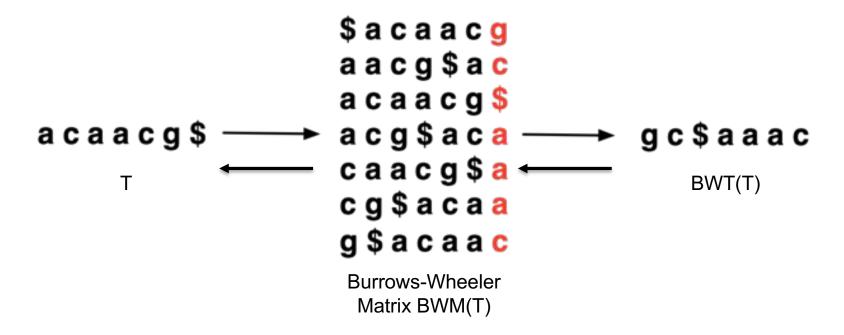
Common to save 50% to 90% on real world files with bzip2

Reversible permutation of the characters in a text



A block sorting lossless data compression algorithm.

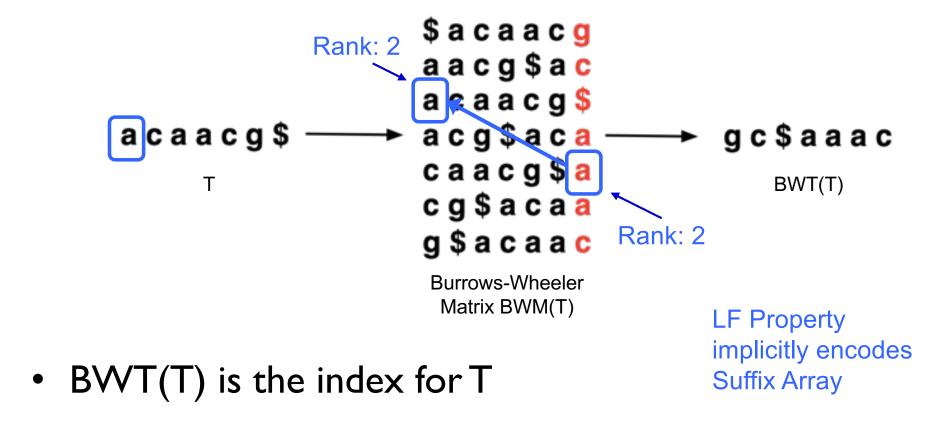
Reversible permutation of the characters in a text



BWT(T) is the index for T

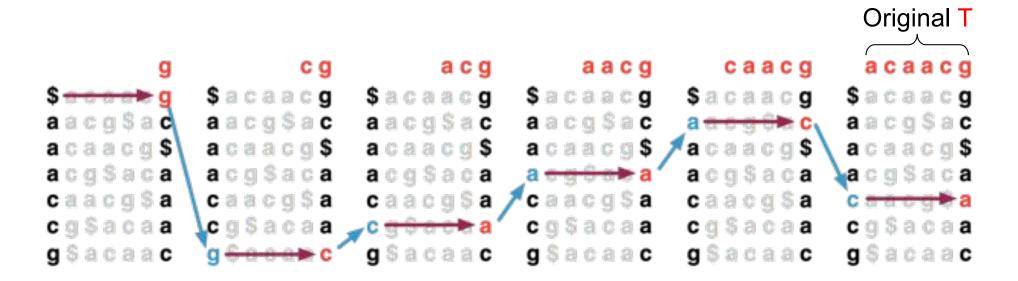
A block sorting lossless data compression algorithm.

Reversible permutation of the characters in a text



A block sorting lossless data compression algorithm.

- Recreating T from BWT(T)
 - Start in the first row and apply LF repeatedly,
 accumulating predecessors along the way



BWT Exact Matching

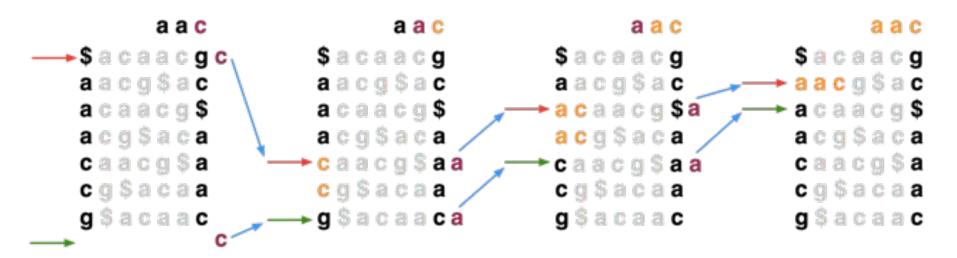
LFc(r, c) does the same thing as LF(r) but it ignores r's actual final character and "pretends" it's c:

```
$acaacg
aacg$ac
acaacg$
acg$aca
caacg$ag
cg$aca
cg$aca
Rank: 2
```

BWT Exact Matching

 Start with a range, (top, bot) encompassing all rows and repeatedly apply LFc:

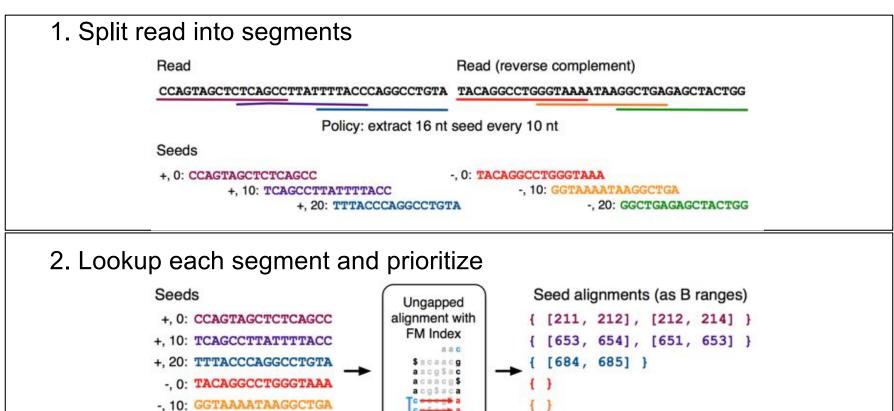
```
top = LFc(top, qc); bot = LFc(bot, qc)
qc = the next character to the left in the query
```



Ferragina P, Manzini G: Opportunistic data structures with applications. FOCS. IEEE Computer Society; 2000.

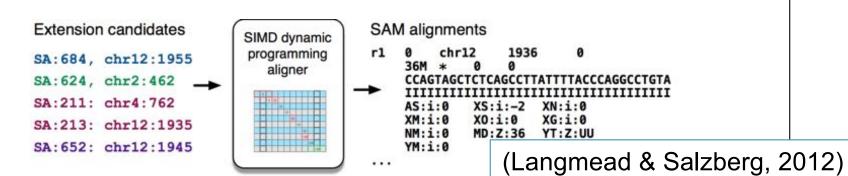
[Search for TTA this BWT string: ACTGA\$TTA]

Algorithm Overview



3. Evaluate end-to-end match

-, 20: GGCTGAGAGCTACTGG



{ [624, 625] }