

Building Large Updatable Colored de Bruijn Graphs via Merging

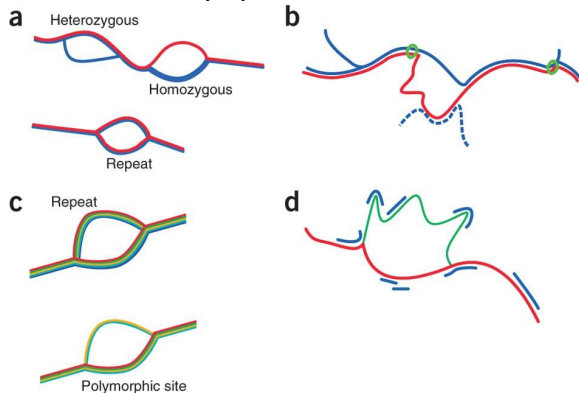
Martin D. Muggli¹ Bahar Alipanahi² , Christina Boucher²

¹Colorado State University

²University of Florida

Colored de Bruijn Graphs

In 2012, Iqbal *et al.* introduced the colored de Bruijn graph with CORTEX. It can detect complex variants within a population without a reference.



Related Work

Efficient de Bruijn graphs

- ABySS
- Conway and Bromage
- Okanohara and Sadakane
- Minia
- BOSS
- MEGAHIT
- Chikhi *et al.*

Efficient colored de Bruijn graphs

- VARI
- Rainbowfish
- Bloom filter trie
- Mantis
- Almodaresi *et al.*

Efficient color representation

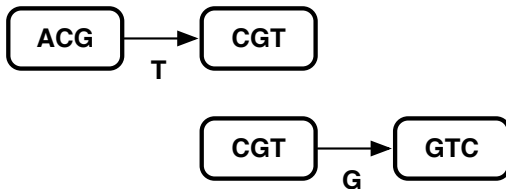
- Mustafa *et al.*
- Mutli-BRWT

Our contribution

- We developed VARI-MERGE
 - Construct succinct colored de Bruijn for sub-populations using VARI
 - New algorithm to merge succinct colored de Bruijn graphs
- Advantages
 - Compress data early \Rightarrow Use less and faster memory
 - Reuse previous work \Rightarrow Incremental update
- First to demonstrate incremental update at this scale

Background: de Bruijn Graphs

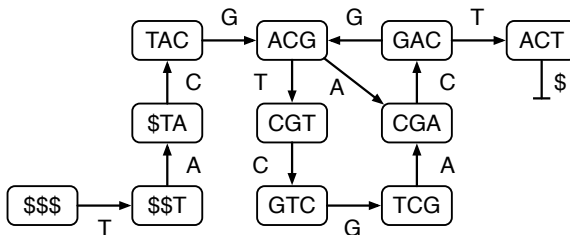
T = TACG**ACG**TCGACT



ACG
CGT

Vertex labels are redundant

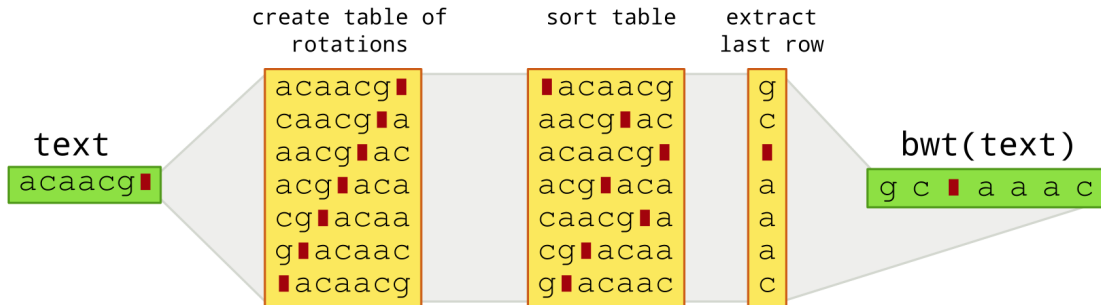
T = TACGACGTCGACT



Burrows-Wheeler Transform (BWT)

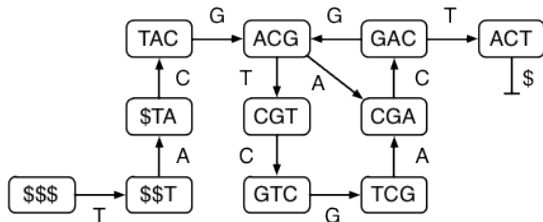
Advantages:

- Compresses repetitive strings well.
- *Self index*: Encodes original string and can provide an index of the implicit *suffix array*.
- $BWT[i] = X[SA[i] - 1]$ if $SA[i] > 1$ and \$ otherwise.



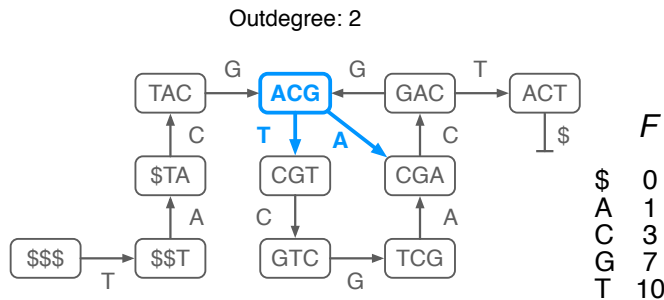
Succinct de Bruijn graphs sort origin labels colex.

Succinct de Bruijn Graphs represent edges as last-to-first mappings in the Burrows-Wheeler transform.



Node			W
\$	\$	\$	T
C	G	A	C
\$	T	A	C
G	A	C	G
G	A	C	T
T	A	C	G
G	T	C	G
A	C	G	A
A	C	G	T
T	C	G	A
\$	\$	T	A
A	C	T	\$
C	G	T	C

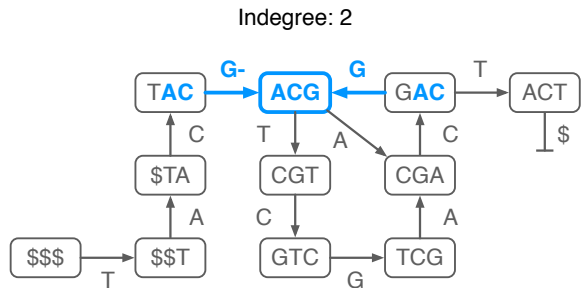
Encoding common origin vertices



	<i>v</i>	<i>i</i>	<i>L</i>	<i>Node</i>	<i>W</i>
	0	0	1	\$ \$ \$	T
	1	1	1	C G A	C
	2	2	1	\$ T A	C
	3	3	0	G A C	G
	4	4	1	G A C	T
	5	5	1	T A C	G-
	6	6	1	G T C	G
	7	7	0	A C G	A
	8	8	1	A C G	T
	9	9	1	T C G	A-
	10	10	1	\$ \$ T	A
	11	11	1	A C T	\$
	12	12	1	C G T	C

	<i>F</i>
\$	0
A	1
C	3
G	7
T	10

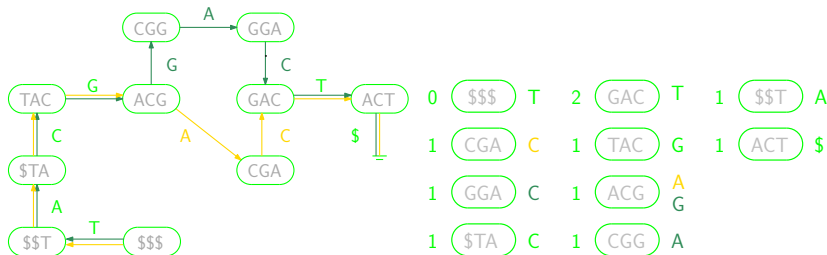
Encoding common destination vertices



	<i>v</i>	<i>i</i>	<i>L</i>	Node	<i>W</i>
	0	0	1	\$ \$ \$	T
	1	1	1	C G A	C
	2	2	1	\$ T A	C
	3	3	0	G A C	G
	4	4	1	G A C	T
	5	5	1	T A C	G-
	6	6	1	G T C	G
	7	7	0	A C G	A
	8	8	1	A C G	T
	9	9	1	T C G	A-
	10	10	1	\$ \$ T	A
	11	11	1	A C T	\$
	12	12	1	C G T	C

	<i>F</i>
\$	0
A	1
C	3
G	7
T	10

VARI method: e.g. a two colored de Bruijn graph and its representation



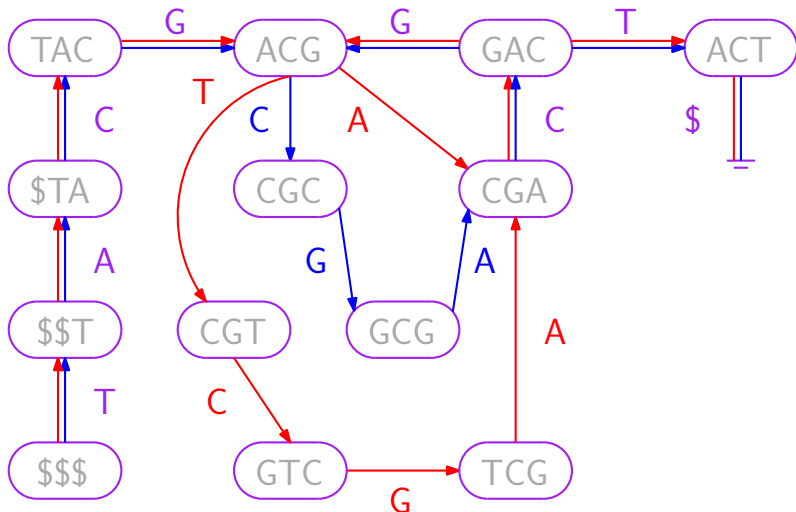
$$\text{EBWT}(G) = \text{TCCCTGAGAA\$}$$

$$C^T = \begin{array}{r} 11011110011 \\ 10111101111 \end{array}$$

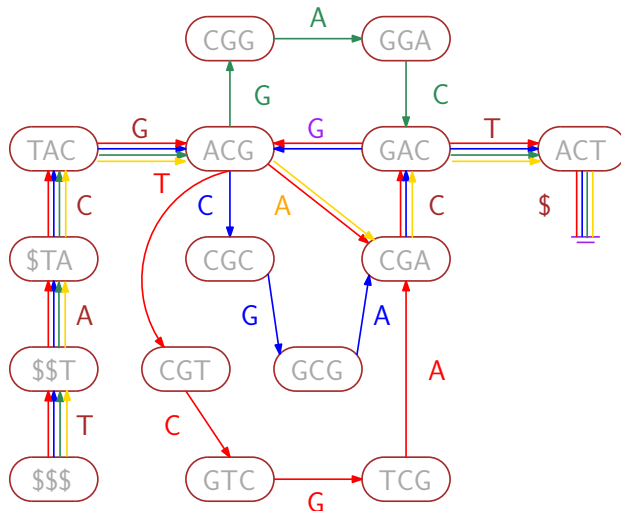
VARI-MERGE: Main Algorithm

- 1 Consider the final population as a collection of sub-populations
- 2 Run VARI on each sub-population
- 3 Run our new algorithm, MERGE on the succinct de Bruijn graphs

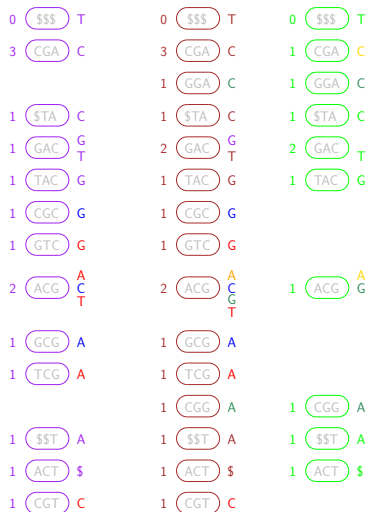
Another two colored de Bruijn graph



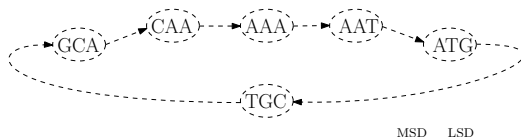
A four colored merged de Bruijn graph



Merging edge labels requires origin vertex label



Two succinct de Bruijn graphs, ignoring color



Edge k -mer 1 A A A T

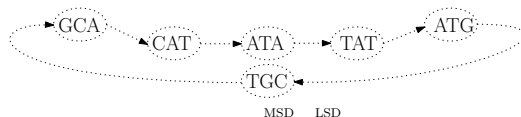
 C A A A

 G C A A

 T G C A

 A T G C

Edge k -mer 6 A A T G



 G C A T

 A T A T

 T G C A

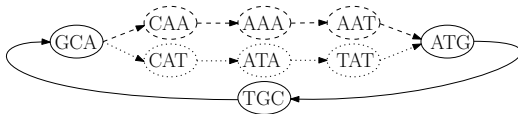
 A T G C

 C A T A

 T A T G

The merged graph

Can we generate the merged succinct graph without reconstructing full vertex k -mers?



Prior BWT merge methods

- Holt and McMillan
- Sirén

Generate Most Significant Digit (MSD) column

	MSD LSD					MSD LSD			
Edge k -mer 1	A	A	A	T		G	C	A	T
	C	A	A	A		A	T	A	T
	G	C	A	A		T	G	C	A
	T	G	C	A		A	T	G	C
	A	T	G	C		C	A	T	A
Edge k -mer 6	A	A	T	G		T	A	T	G

Generate common suffix groups for vertex labels

	MSD	LSD	
Edge k -mer 1	A	A	A
	C	A	A
	G	C	A
	T	G	C
	A	T	G
Edge k -mer 6	A	A	T

	MSD	LSD	
	G	C	A
	A	T	A
	T	G	C
	A	T	G
	C	A	T
	T	A	T

Generate next most significant digit column

		MSD	LSD	
Edge k -mer 1	A	A	A	T
	C	A	A	A
	G	C	A	A
	T	G	C	A
	A	T	G	C
Edge k -mer 6	A	A	T	G

	MSD	LSD	
G	C	A	T
A	T	A	T
T	G	C	A
A	T	G	C
C	A	T	A
T	A	T	G

Recursively subdivide existing groups using new column

	MSD	LSD	
Edge k -mer 1	A	A	A
	C	A	A
	G	C	A
	T	G	C
	A	T	G
Edge k -mer 6	A	A	T

	MSD	LSD	
	G	C	A
	A	T	A
	T	G	C
	A	T	G
	C	A	T
	T	A	T

Constructing 8,000 Salmonella sample graph via merging

Program	Time	External Memory	RAM
VARI	37 h 27 min	4.6 TB	271 GB
VARIMERGE	26 h 30 min	1.5 TB	137 GB

Incremental update performance

Program	Time	External Memory	RAM
VARI(16000)	69 h 8 min	2.34 TB	254 GB
VARI(1)	7 s	460 MB	2.3 GB
MERGE(16000, 1)	7 h 9 min	0	254 GB

Comparison of space-efficient colored graph construction methods

Dataset	No. of k -mers	Program	Output Size	Time	RAM(RSS)
16,000	5.8 Billion	VARI / Rainbowfish	N/A	N/A	N/A
		Bloom Filter Trie	N/A	N/A	N/A
		Multi-BRWT	N/A	N/A	N/A
		Mantis / Method of Almodaresi et al.	256 GB	36 h 12 m	316 GB
		VARIMERGE	233 GB	69 h 8 m	254 GB

Conclusion

- Uncompressed work in small chunk reduces external memory
- Reusing previous computational work lets us build an updated version
- Radix based method satisfies metadata consistency and has no random access

Future work

- What is the optimal sub-population size for initial succinct colored de Bruijn graphs?
- VARIMERGE is radix based, would a trie based merge like bwt-merge be superior under some circumstances?

Acknowledgements

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National Institute of
Allergy and
Infectious Diseases

Questions

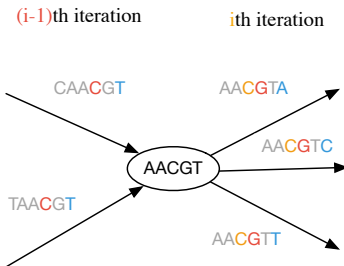
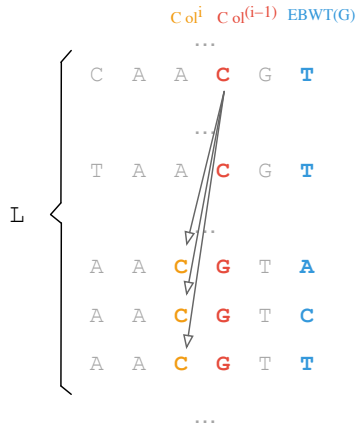
Questions?

Validation

- 1 Build succinct colored de Bruijn graphs for *E. coli* genome datasets A and B independently using VARI
- 2 Merge them using VARIMERGE
- 3 Build a succinct colored de Bruijn graph for all the data in A and B using VARI
- 4 Compare merged graph from step 2 with directly constructed graph from step 3

Result: bit-for-bit identical files on disk

Efficiently computing one column at a time



Constructing 8,000 color graph via merging

	Input Stats		de Bruijn Graph			Color Matrix			Combined Requirements			
Program and Dataset	k-mers	Colors	RAM	Time	Size	RAM	Time	Size	RAM	Ext. Mem.	Time	Size
VARI(4A)	1.1 B	4,000	136 GB	8 h 46 m	0.31 GB	52 GB	1 h 39 m	51.2 GB	136 GB	1 TB	10 h 25 m	51 GB
VARI(4B)	1.5 B	4,000	137 GB	10 h 40 m	0.52 GB	54 GB	2 h 22 m	52.5 GB	137 GB	1.5 TB	13 h 2 m	53 GB
MERGE(4A, 4B)	2.4 B	8,000	10 GB	2 h 1 m	0.63 GB	117 GB	1 h 2 m	106 GB	117 GB	0 TB	3 h 3 m	106 GB
VARI-MERGE	2.4	8,000	137 GB	21 h 27 m	0.63 GB	117 GB	5 h 3 m	117 GB	137 GB	1.5 TB	26 h 30 m	106 GB

Constructing 16,000 color graph via merging

Program and Dataset	Input Stats		de Bruijn Graph			Color Matrix			Combined Requirements			
	<i>k</i> -mers	Colors	RAM	Time	Size	RAM	Time	Size	RAM	Ext. Mem.	Time	Size
VARI(4C)	1.7 B	4,000	135 GB	10 h 53 m	0.46 GB	53 GB	2 h 34 m	51.8 GB	135 GB	1.6 TB	13 h 27 m	52 GB
VARI(4D)	2.4 B	4,000	137 GB	14 h 35 m	0.67 GB	59 GB	3 h 37 m	57.9 GB	137 GB	2.34 TB	18 h 12 m	59 GB
MERGE(4C, 4D)	3.8 B	8,000	17 GB	2 h 59 m	1.00 GB	118 GB	57 m	107 GB	118 GB	0 TB	3 h 56 m	108 GB
MERGE(8AB, 8CD)	5.8 B	16,000	25 GB	4 h 53 m	1.60 GB	254 GB	2 h 10 m	232 GB	254 GB	0 TB	7 h 3 m	233 GB
VARI-MERGE	5.8 B	16,000	137 GB	54 h 47 m	1.60 GB	254 GB	14 h 21 m	232 GB	254 GB	2.34 TB	69 h 8 m	233 GB

FM-Index and Backward Search

Advantages:

- Exact search in $O(n)$.
- Compressed Suffix Array (CSA)

