Constructing the CDAWG CFG using LCP-Intervals

Alan Cleary and Jordan Dood

Some Definitions

CFG - <u>C</u>ontext <u>Free</u> <u>G</u>rammar

CDAWG - <u>Compacted Directed Acyclic Word Graph</u>

LCP - <u>L</u>ongest <u>C</u>ommon <u>P</u>refix

Context Free Grammars

(CFGs)

- A CFG is a set of production rules
- Recursing the rules from a designated 'start' rule yields a string or language of strings
- Grammars that produce a single string are called 'straight-line' grammars

Why We Are Interested in CFGs

- CFGs can be built from repeated sub-sequences, and thus can result in compression, of the string
- CFGs are self-indexing
- CFGs can be constructed in a number of different ways, which offers flexibility



Motivation

What is needed

We wanted to build a 'bridge' between the CFG realm and that of other well studied string data structures

Putting It Another Way

We want to find a way of compressing strings as a context free grammars that

maintains the information structure related to other data structures, like suffix trees, CDAWGS, ect

and ...

does so efficiently in both time and space, ideally O(n) time where *n* is the size of the string

Design Criteria

1. Compress data into a straight-line

CFG

2. Build that CFG using the maximal

repeats of the string

3. With a time complexity of

O(nlogn) or O(n)

$S_{SA[i]}$	1	LCP-intervals			
\$ AGAGCGAGAGCGCGC\$	0	2	6	ľ	
AGCGAGAGCGCGC\$			4		
C\$		1		l r	
CGAGAGCGCGC5 CGC\$			2	3	
GAGAGCGCGC\$		1	3		
GAGCGAGAGCGCGC\$ GAGCGCGCC\$				5	
GC\$ GCGAGAGCGCGC\$			2		
GCGC\$ GCGCGC\$				3 4	
	-	_			

Repeats

A repeat is substring that occurs more than once in a string

A <u>right-maximal repeat</u> is repeat that cannot be extended further to the right without reducing the number of occurrences

A <u>left-maximal repeat</u> is is a repeat that cannot be extended further to the left without reducing the number of occurrences

A <u>maximal repeat</u> is a repeat that is both right- and left-maximal and is a subset of both sets

Repeats

These repeats are inherent to a given string

Right-maximal repeats are represented by the internal nodes of both the suffix tree and the CDAWG, and define the <u>L</u>ongest <u>Common Prefix (LCP) intervals of the suffix</u> array, the intervals of a suffix array that share a common prefix

Because of this connection to these data structures we wanted to utilize maximal repeats to build our grammar

$S_{SA[i]}$	1	LCP-intervals				
\$ AGAGCGAGAGCGCGC\$			6			
AGAGCGCGC\$ AGCGAGAGAGCGCGC\$	0	2	4			
AGCGCGCS C\$		1]		
CGAGAGCGCGC\$ CGC\$			2	3		
CGCGC\$ GAGAGCGCGC\$		1				
GAGCGAGAGCGCGC\$ GAGCGCGC\$			3	5		
GC\$ GCGAGAGCGCGC\$			2			
GCGC\$ GCGCGC\$				3 4		

The Crux



... AACTGGTCGATCGATCGATCTAGGCTACATGGCTAGCCATCTACTGCTGACTGGATCGACTAGAA ...



The difficulty is in choosing which rules to include and how to handle overlaps



... AACTGGTCGATCGATCGATCTAGGCTACATGGCTAGCCATCTACTGCTGACTGGATCGACTAGAA ...

Strait-line Grammar from Suffix Tree

This process uses a <u>Compacted</u> <u>Directed</u> <u>Acyclic</u> <u>Word</u> <u>Graph</u> (CDAWG)

As well as a number of simplification steps, to build a CFG from the structure of the CDAWG



Belazzougui, Djamal, and Fabio Cunial. "Representing the suffix tree with the CDAWG." 28th Annual Symposium on Combinatorial Pattern Matching (2017).

Key Inspiration

For CDAWG-CFG

Because there is a bijection between the internal nodes of the suffix tree and the right-maximal repeats, and because a subset of these nodes are preserved during the transformations, the grammar that results is comprised of these repeats.

We exploit this by noting the bijection between the internal nodes of a CDAWG and maximal repeats.

Why This Matters to Us

• Maximal repeats are (by definition) a subset of the right-maximal repeats, thus maintaining the relatedness to other data structures

• Since the Belazzougui, et al. work shows that right-maximal repeats can be used to build a grammar, and crucially, how these repeats are used to encode the target string, maximal repeats should be able to do something similar

• The downside is that this starts with a suffix tree and has a lot of intermediate graph structures

Comparison of Algorithms

Belazzougui, et al.

- Requires precomputting a suffix tree and CDAWG
- Uses CDAWG nodes (right-maximal repeats)
- Uses extraneous (for our purpose) steps

Our Algorithm

• Uses LCP-intervals, which can

be computed from many data

structures

- Can be computed online
- Computes the CFG directly

Our Approach

String





Start

CFG



Our data structure is an interval tree for answering stabbing queries on LCP-intervals in constant time, or more generally nested intervals in a finite integer range



<- Pointer Map



<- Pointer Map
<- Bit Vector





https://www.shutterstock.com/search/dagger-silhouette

2-Modes of Action

Our data structure can be built first, then preprocessed in O(n)so that 'stabbing' is done in O(1)

Or it can be used <u>online</u> which allows for stabbing queries to be done in *O(log n)* time (binary search)

This duality is possible because of the nested structure of LCP-intervals and the order of their traversal

Preprocessed O(n) time



Online O(n log n) time

Iterate LCP-intervals, Building Data Structure and Encoding Grammar



Algorithm

(Optimal)

1) Iterate the LCP-intervals



Beller, Timo, Katharina Berger, and Enno Ohlebusch. "Space-efficient computation of maximal and supermaximal repeats in genome sequences." *International Symposium on String Processing and Information Retrieval*. Springer, Berlin, Heidelberg, 2012.

1) Iterate the LCP-intervals



The iteration is done in first length, then lexicographic order

Beller, Timo, Katharina Berger, and Enno Ohlebusch. "Space-efficient computation of maximal and supermaximal repeats in genome sequences." *International Symposium on String Processing and Information Retrieval*. Springer, Berlin, Heidelberg, 2012.

1) Iterate the LCP-intervals



<u>NOTE:</u> LCP-intervals can be computed from a variety of string data structures, this one uses an FM-index

Beller, Timo, Katharina Berger, and Enno Ohlebusch. "Space-efficient computation of maximal and supermaximal repeats in genome sequences." *International Symposium on String Processing and Information Retrieval*. Springer, Berlin, Heidelberg, 2012.

2) Determine Maximality



3) Add to Stabbing Data Structure



4) Repeat (Steps 1-3)





. . .

5) Process The Stabbing Data Structure

AATCCTCATCGTCCATG

••••



6) Encode The Rules

AATCCTCATCGTCCATG

•••



6) Encode The Rules

AATCCTCATCGTCCATG



. . .

7) Repeat (step 6)



8) Encode The Start Rule

AATCCTCATCGTCCATG

String



...



Our Implementation

https://github.com/alancleary/cdawg-cfg

(and some results)

Conclusions

Our Algorithm . . .

• Compresses data into a CFG that is based on maximal repeats

• This can be done online or not, with time complexity of *O*(*n* log *n*) or *O*(*n*), respectively

• We achieve this using a novel interval stabbing data structure to reduce intermediate steps and data structures

Future Work

• Use the relation to other string data structures to port over functionality

• Better characterize and improve the compression ratio for our grammars

• Explore the opportunities for operating on compressed data

Funding and Acknowledgements



This work was supported by NSF Award Number 2105391

National Center for Genome Resources

