

Minimal Absent Words

Alice Heliou

October 13, 2015

- 1 Introduction
- 2 Minimal Absent Words
- 3 Computation of Minimal Absent Words

Outline

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'Negative' information

Principle

Given a sequence of letters, we focus on words that don't occur. Their absence may have a signification.

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Example

In a random sequence S , we expect that every word of size less than $\log_\sigma(|S|)$ occurs in S .

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Example

In a random sequence S , we expect that every word of size less than $\log_{\sigma}(|S|)$ occurs in S .

The human genome contains around 3G nucleotides (A, C, G, T). Yet some words of size 11, are absent ($11 < \log_4(3 * 10^9) = 15,7$)

'Negative' information

Application

Three minimal sequences found in Ebola virus genomes and absent from human DNA, [Silva et al.], 2015

3 small sequences (of length between 12 and 14) that appear in the Ebola genome as coding for proteins, are absent from the Human genome.

This was done by analyzing 99 virus and the Human genome reference GRC-38.

Sequence

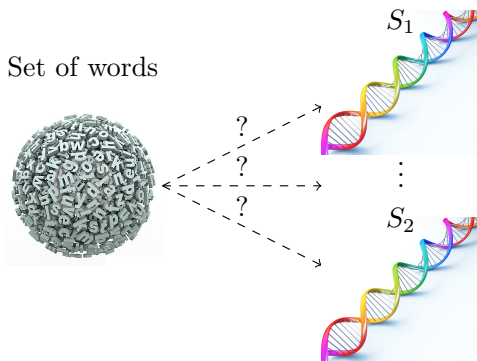


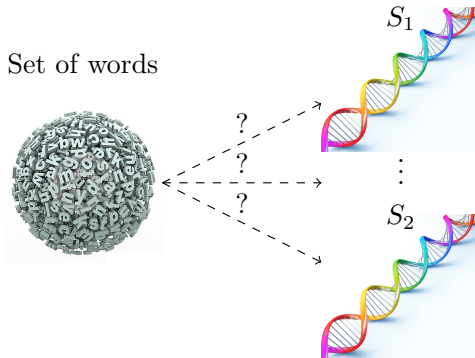
Unicity



Set of Absent Words







Property

For each set of words \mathcal{M} if there exists a sequence \mathcal{S} such that \mathcal{M} is its set of absent words, then \mathcal{S} is unique.

Absent words are too numerous

The number of absent words from a sequence of size n is **exponential** in n .

There are at most two words of size $n - 1$ that occur in $S \Rightarrow$ at least $\sigma^{n-1} - 2$ absent words of size $n - 1$
with σ the size of the alphabet.

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Definition : Minimal Absent Word

A minimal absent word of a sequence is an absent word whose proper factors (longest prefix, and longest suffix) all occur in the sequence.

An upper bound on the number of minimal absent words is $\mathcal{O}(\sigma n)$.

Crochemore et al. 1998, Mignosi et al. 2002

$$\begin{array}{cccccccc} & 0 & 1 & 2 & 3 & 4 & 5 & 6 & 7 \\ S = & A & A & C & A & C & A & C & C \end{array}$$

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$$\begin{array}{cccccccc}
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 S = & \textcolor{red}{A} & \textcolor{red}{A} & C & A & C & A & C & C
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An absent word has a minimal absent word as factor

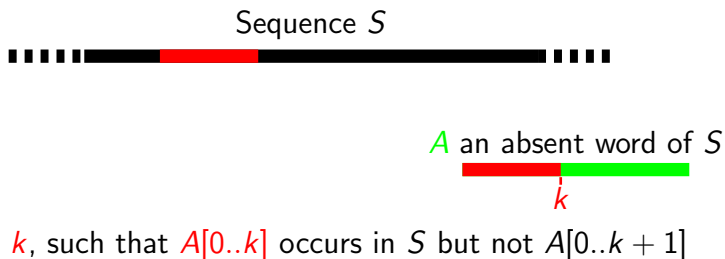
Sequence S



A an absent word of S



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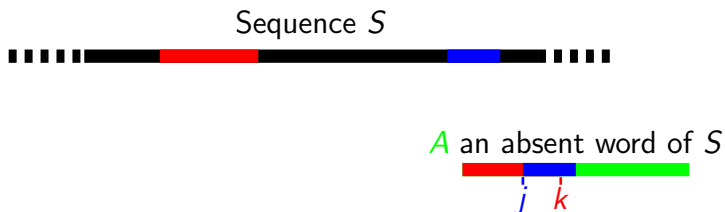
A an absent word of S



k , such that $A[0..k]$ occurs in S but not $A[0..k+1]$

j , such that $A[j..k+1]$ occurs in S but not $A[j-1..k+1]$

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$A[j-1..k+1]$ is a minimal absent word of S

because $A[j..k+1]$ and $A[j-1..k]$ occur in S .

Retrieving the sequence from its set of minimal absent words

Retrieving a sequence from its set of minimal absent words can be done in linear time \Rightarrow Gabriele Fici thesis Minimal Forbidden Words and Applications (2006).

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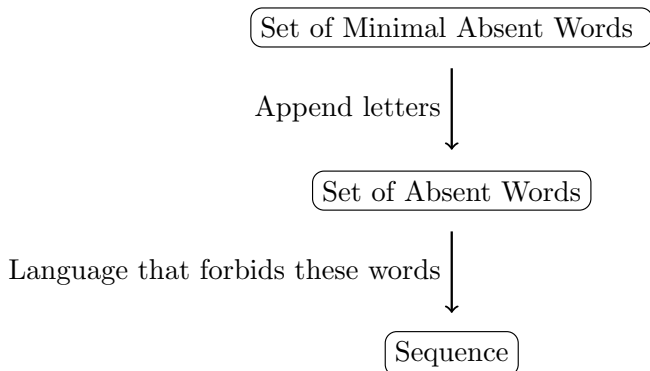
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Why can we retrieve it ?

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Definition: Maximal repeated pair

A maximal repeated pair in a S is a triple (i, j, w) such that:

- w occurs in S at positions i and j
- $S[i - 1] \neq S[j - 1]$
- $S[i + |w|] \neq S[j + |w|]$

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longest prefix of A

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longest suffix of A

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How to find all maximal repeated pairs ?

By sorting all the suffixes of S , \Rightarrow Suffix Array

Suffix Array by Manber& Myers in 1990

Table containing the starting position of the suffixes when they are in alphabetical order. It allows fast localisation of patterns.

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$$\begin{array}{cccccccc}
 0 & 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 \\
 S = & A & A & C & A & C & A & C & C \#
 \end{array}$$

0	A	A	C	A	C	A	C	C	#
1	A	C	A	C	A	C	C	#	
2	C	A	C	A	C	C	#		
3	A	C	A	C	C	#			
4	C	A	C	C	#				
5	A	C	C	#					
6	C	C	#						
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8	#								

Suffixes of y

Ordered suffixes of S

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	<i>pos</i>										
0	A	A	C	A	C	A	C	C	#	8	#
1	A	C	A	C	A	C	C	#			
2	C	A	C	A	C	C	#				
3	A	C	A	C	C	#					
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 $S = \text{AACACACC}\#$

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0	A A C A C A C C #	8	#
1	A C A C A C C #	0	A A C A C A C C #
2	C A C A C C #		
3	A C A C C #		
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4	C A C C #	5 A C C #
5	A C C #	7 C #
6	C C #	2 C A C A C C #
7	C #	4 C A C C #
8	#	6 C C #

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8	#								

Suffixes of y

\Rightarrow

SA	
8	#
0	A A C A C A C C #
1	A C A C A C C #
3	A C A C C #
5	A C C #
7	C #
2	C A C A C C #
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6	C C #

Ordered suffixes of S

Computation of minimal absent words

Pre-computation

Construction :

- Suffix Array, linear time and space since 2003
- Longest Common Prefix table, linear time and space with the SA and the sequence as input

Computation

- Travel twice through those tables, in order to construct the set of letters that occurs just before each right-maximal repetition.
- Deduce the set of minimal absent words.

Applications

Biology

- Linear-Time Sequence Comparison Using Minimal Absent Words & Applications, [Crochemore et al.], 2015
- Minimal Absent Words in Prokaryotic and Eukaryotic Genomes, [Garcia et al.], 2011

Computer Science

- Data Compression Using Antidictionaries, [Crochemore et al.], 2000, [Fiala and Holub], 2008

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Before 2014

References	Time for fixed size alphabet	Space	Structure
Crochemore et al. 1998 Automata and forbidden words	$\mathcal{O}(n)$	$\mathcal{O}(n)$	suffix automata
Pinho et al. 2009 On finding minimal absent words	$\mathcal{O}(n^2)$	$\mathcal{O}(n)$	suffix array
Belazzougui et al. 2013 Versatile Succinct Representations of the Bidirectional Burrows-Wheeler Transform.	$\mathcal{O}(n)$	$\mathcal{O}(n)$	compact bidirectional BWT

Constants reduction

References	Time for fixed size alphabet	Space	Structure
Ota et al. 2014 Dynamic construction of an antidictionary with linear complexity Theoretical Computer Science	$\mathcal{O}(n)$	$\mathcal{O}(n)$	suffix tree, dynamic approach
Barton et al. 2014 Linear-time computation of minimal absent words BMC Bioinfo	$\mathcal{O}(n)$	$\mathcal{O}(n)$	suffix array
Belazzougui et al. 2015 Space-efficient detection of unusual words. CPM	randomized $\mathcal{O}(n)$	$\mathcal{O}(n)$	BWT & few additional structures
Barton et al. 2015 Engineering the Computation of Minimal Absent Words. PPAM	$\mathcal{O}(n/p)$	$\mathcal{O}(n)$	suffix array

Perspectives

- Use external memory computation to find a trade-off between running time and RAM usage.
- Knowing the set of minimal absent words of a sequence, deduce the set of a circular shift.

Thank you

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Questions ?

Linear time construction of the suffix array

over integer alphabets

The skew algorithm, Karkkainen and Sanders 2003

Main idea : Divide suffixes into 2 groups :

- Those starting a position $i \not\equiv 0 \pmod 3$
- Those starting a position $i \equiv 0 \pmod 3$

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$$\begin{array}{cccccccccccccc}
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 S = & M & I & S & S & I & S & S & I & P & P & I & \# \#
 \end{array}$$

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Outline :

- recursively sort the suffixes of the first group
- merge with the second group

first step, recursively sort the suffixes starting at $i \not\equiv 0 \pmod 3$

- Consider all the triples of starting at positions $i \not\equiv 0 \pmod 3$.

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- Assign them lexicographical names $\Rightarrow S'$
This can be done in linear time by radix sort and recursion
 - if some of them get the same lexicographic name, we compute recursively the suffix array of the string S^{12}

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This can be done in linear time by radix sort and recursion
 - if some of them get the same lexicographic name, we compute recursively the suffix array of the string S^{12}
- Once all the triples are ordered, we have the ordering of the suffixes starting at $i \not\equiv 0 \pmod 3$

handling 0 suffixes

- Sort the group 0 suffixes, using the representation $(S[i], S_{i+1})$

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 - if $i \equiv 1 \pmod 3$, we compare the pair representations $(s[j], S_{j+1})$ and $(s[i], S_{i+1})$
 - if $i \equiv 2 \pmod 3$, we compare the triple representations $(s[j], s[j+1], S_{j+2})$ and $(s[i], s[i+1], S_{i+2})$