

How many double squares can a string contain?

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Outline

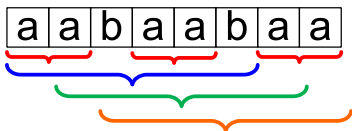
- 1 Motivation and background
- 2 Basic notions and tools
- 3 Double squares
- 4 Inversion factors
- 5 Rightmost double squares
- 6 An upper bound for the number of double squares
- 7 Main theorems
- 8 Conclusion

Motivation and background

We are dealing with finite strings over finite alphabets. There is no particular requirement about the order of the alphabet.

What is the *maximum number of distinct squares problem* ?

We are counting types of squares rather than their occurrences.

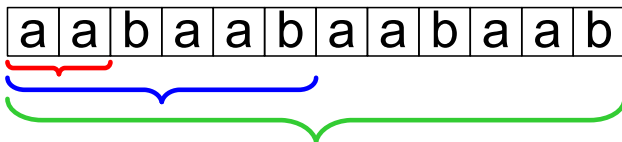


has 6 occurrences of squares, but only 4 distinct squares, *aa*, *aabaab*, *abaaba*, and *baabaa*.

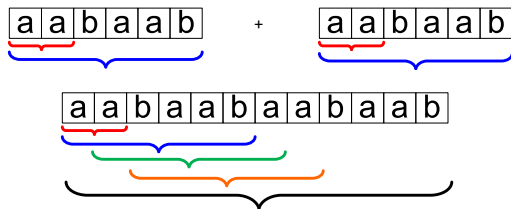
A trivial bound: the number of all occurrences of primitively rooted squares in a string of length n is bounded by $O(n \log n)$ (*Crochemore 1978*) and the number of distinct non-primitively rooted squares is $O(n)$ (*Kubica et al. 2013*)

Could it be $O(n)$? And if so, what would be the constant?

Why this is not simple? In a string of length n , $O(\log n)$ squares can start at the same position!



It is easy to compute it for short strings, so why induction cannot be used?



Concatenation does both “destroys” existing types through multiple-occurrences and “creates” new types. Of course, same holds true for the reverse process - partitioning of strings.

Theorem (Fraenkel-Simpson, 1998)

There are at most $2n$ distinct squares in a string of length n .

Count only the *rightmost* occurrences. Fraenkel-Simpson showed that if there are three rightmost squares uu , vv , and ww starting at the same position so that $|u| < |v| < |w|$, then ww contains a farther copy of uu , based on Crochemore-Rytter (1995) Lemma showing that in such a case, $|w| \geq |u| + |v|$.

Fraenkel-Simpson hypothesized that the number of distinct squares should be bounded by n , i.e.

$$\sigma(n) \leq n$$

where $\sigma(n) = \max \{ s(x) : x \text{ is a string of length } n \}$.

Fraenkel-Simpson gave an infinite sequence of strings $\{x_n\}_{n=1}^{\infty}$ so that $|x_n| \nearrow \infty$ and

$$\frac{s(x_n)}{|x_n|} \nearrow 1$$

where $s(x) = \text{number of distinct squares in } x$.

- In 2005 *Ilie* provided a simpler proof of *Fraenkel-Simpson's* Theorem and in 2007 presented an asymptotic upper bound of $2n - \theta(\log n)$.
- In 2011 *Deza-F.* proposed a d -step approach to the problem and conjectured that $\sigma_d(n) \leq n - d$, where $\sigma_d(n) = \max \{ s(x) : x \text{ is a string of length } n \text{ with } d \text{ distinct symbols} \}$.

Basic notions and tools

Definition

non-trivial power of a string x is a concatenation of m copies of x denoted as x^m ; x^2 is a *square*, x^3 a *cube*.

A string x is *primitive* if $x \neq y^n$ for any y and any $n \geq 2$.

primitive root of x is the shortest y so that $x = y^n$.
(Note that y must be primitive.)

x and y are *conjugates* if $x = uv$ and $y = vu$ for some u, v .

Lemma (*Synchronization principle*)

Given a primitive string x , a proper suffix y of x , a proper prefix z of x , and $m \geq 0$, there are exactly m occurrences of x in yx^mz .



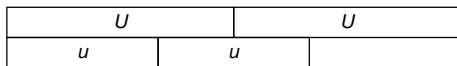
Lemma (*Common factor lemma*)

For any strings x and y , if a non-trivial power of x and a non-trivial power of y have a common factor of length $|x|+|y|$, then the primitive roots of x and y are conjugates.

Double squares

- *Fraenkel-Simpson*: only two rightmost squares can start at the same position. Thus, only one rightmost square or two rightmost squares may start at any position.
- *Lam (2009 – unpublished)* tried bounding the number of *double squares* and hence bound the number of distinct squares. His approach is based on a taxonomy of all possible configurations of two double squares yielding a bound of $\frac{94}{48}n \approx 1.98n$.

A configuration of two squares



has been investigated in many different contexts:

- *Smyth et. al.*: with intention to find a position for amortization argument for runs conjecture.
- in computational framework by *Deza-F.-Jiang*: such configurations are used in *Liu's* Ph.D. thesis to speed up computation of $\sigma_d(n)$.
- *Lam*: two rightmost squares have a unique structure.

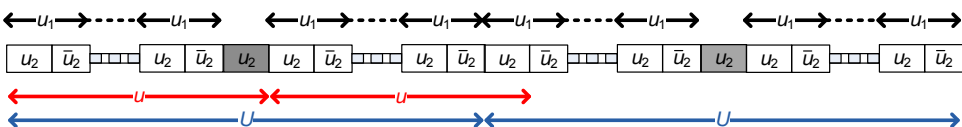
Lemma

Let uu and UU be two squares in a string x starting at the same position with $|u| < |U|$ such that either

- (a) both uu and UU are rightmost occurrences, or*
- (b) uu or UU is primitively rooted and $|U| < |uu|$*

Then $|u| < |U| < |uu| < |UU|$ and there is a unique primitive string u_1 , a unique proper prefix u_2 of u_1 , and unique integers e_1 and e_2 satisfying $1 \leq e_2 \leq e_1$ such that $\mathbf{u} = \mathbf{u}_1^{e_1} \mathbf{u}_2$ and $\mathbf{U} = \mathbf{u}_1^{e_1} \mathbf{u}_2 \mathbf{u}_1^{e_2}$; i.e. uu and UU form a double square.

$$u_1^{u(1)} u_2 u_1^{u(2)} u_1^{u(1)} u_2 u_1^{u(2)}$$



Thus, only strings of length at least 10 may contain a double square: $|UU| = 2((u(1)+u(2))|u_1|+|u_2|) \geq 2((1+1)2+1) = 10$.

Cyclic shift (rotation) to the right is controlled by

$$lcp(u_1, \bar{u}_1)$$

while cyclic shift to the left is controlled by

$$lcs(u_1, \bar{u}_1)$$

lcp = largest common prefix

lcs = largest common suffix

$$u_1 = aabaa, u_2 = aab, \bar{u}_2 = aa, u(1) = u(2) = 2$$

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$u_1 = aabaa$, $u_2 = aab$, $\bar{u}_2 = aa$, $u(1) = 2$, and $u(2) = 1$.

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Definition

For a double square \mathcal{U} , $\bar{v}vv\bar{v}$ where $|\bar{v}| = |\bar{u}_2|$ and $|v| = |u_2|$ is an *inversion factor*

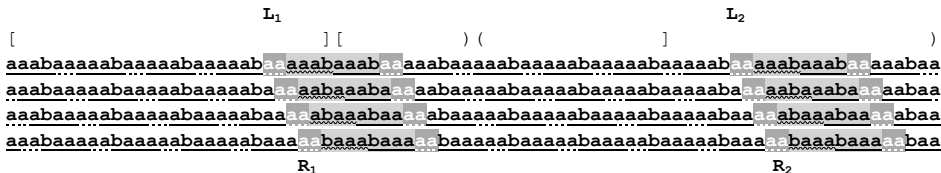
$$\mathcal{U} = u_1^{u(1)} u_2 u_1^{u(2)+u(1)} u_2 u_1^{u(2)} =$$

$$u_1^{(u(1)-1)} u_2 \bar{u}_2 u_2 u_2 \bar{u}_2 u_1^{u(2)+u(1)-2} u_2 \bar{u}_2 u_2 u_2 \bar{u}_2 u_1^{(u(2)-1)}$$

 N_1
 N_2

natural inversion factors

A cyclic shift of an inversion factor is an inversion factor, also controlled by $lcp(u_1, \bar{u}_1)$ and $lcs(u_1, \bar{u}_1)$.



All inversion factors are cyclic shifts of the natural ones:

Lemma (*Inversion factor lemma*)

Given a double square \mathcal{U} , there is an inversion factor of \mathcal{U} within the string UU starting at position $i \iff i \in [L_1, R_1] \cup [L_2, R_2]$.

Inversion factor lemma for distinct squares

Theorem (*Fraenkel-Simpson, Ilie*)

At most two rightmost squares can start at the same position.

Let us assume that 3 rightmost squares uu , UU , vv start at the same position.

By item (c) of Inversion factor lemma, uu and UU form a double square \mathcal{U} : $u = u_1 \mathcal{U}^{(1)} u_2$ and $U = u_1 \mathcal{U}^{(1)} u_2 u_1 \mathcal{U}^{(2)}$.

Since the first v contains an inversion factor, the second v must also contain an inversion factor.

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If the inversion factor in the second v were from $[L_2, R_2]$, then $|v| = |U|$, a contradiction.

Hence v must not contain an inversion factor from $[L_2, R_2]$ and so $u_1^{u(1)}u_2u_1^{u(1)+u(2)-1}u_2$ must be a prefix of v .

Therefore vv contains another copy of $u_1^{u(1)}u_2u_1^{u(1)}u_2 = uu$, a contradiction.

Fundamental Lemma:

Lemma

Let x be a string starting with a double square \mathcal{U} . Let \mathcal{V} be a double square with $s(\mathcal{U}) < s(\mathcal{V})$, then either

(a) $s(\mathcal{V}) < R_1(\mathcal{U})$, in which case either

(a₁) \mathcal{V} is an α -mate of \mathcal{U} (cyclic shift), or

(a₂) \mathcal{V} is a β -mate of \mathcal{U} (cyclic shift of U to V), or

(a₃) \mathcal{V} is a γ -mate of \mathcal{U} (cyclic shift of U to v), or

(a₄) \mathcal{V} is a δ -mate of \mathcal{U} (big tail),

or

(b) $R_1(\mathcal{U}) \leq s(\mathcal{V})$, then

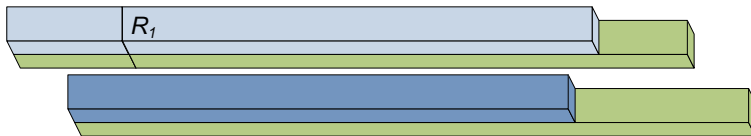
(b₁) \mathcal{V} is a ε -mate of \mathcal{U} (big gap).

α -mate (cyclic shift):



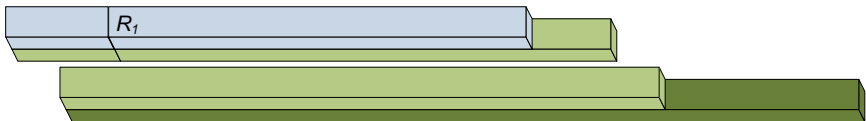
[[] [] ([])]
aaabaaaaab aaaaabaaab aaaaabaaaaab aaaaabaaab aa . . .
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β -mate (cyclic shift of U to V)



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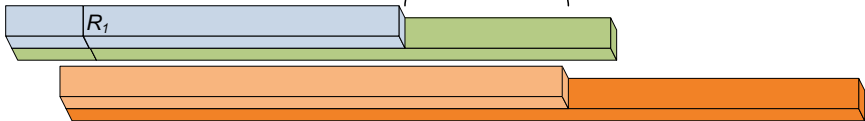
γ -mate (cyclic shift of U to v)



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δ -mate (big tail)

sufficiently big tail

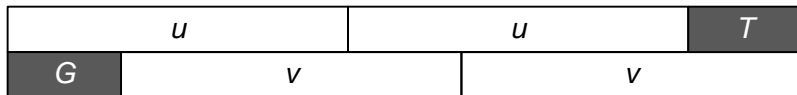


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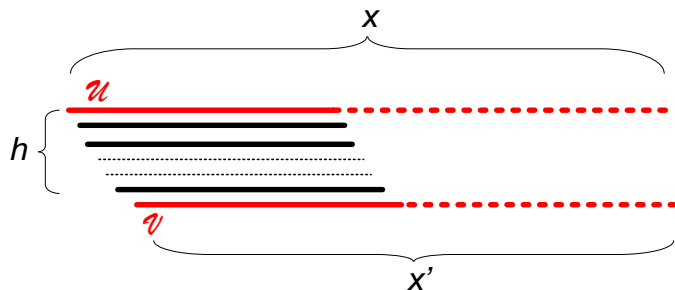
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An upper bound for the number of double squares

We show by induction a bound $\delta(x) \leq \frac{5}{6}|x| - \frac{1}{3}|u|$, where uu is the shorter square of the leftmost double square of x .



The fundamental lemma basically says that either the **gap** $G(u, v)$ is “big” or the **tail** $T(u, v)$ is “big” (for δ -mate and ε -mate), or it is case of α -mate, β -mate, or γ -mate.



Lemma (Gap-Tail lemma)

$\delta(x') \leq \frac{5}{6}|x'| - \frac{1}{3}|v|$ implies

$$\delta(x) \leq \frac{5}{6}|x| - \frac{1}{3}|u| + h - \frac{1}{2}|G(u, v)| - \frac{1}{3}|T(u, v)|$$

We deal with α -mates, β -mates, and γ -mates separately.

It is possible as they form families, either a pure α -family, or $\alpha+\beta$ -family, or $\alpha+\beta+\gamma$ -family.

\mathcal{U} -family consists only of α -mates

Illustration of α -family with $u(1) = u(2)$

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It is easy to bound the size of α -family, as it is controlled by $lcp(u_1, \bar{u}_1)$ and $lcp(y, u_2)$ where y is x without UU : the size $\leq |u_1|$.

- Either there are no other double squares, and then it can be shown directly that the bound holds, or
- There is a \mathcal{V} underneath, and we can use induction using the Gap-Tail lemma. \mathcal{V} must be either γ -mate, or δ -mate, or ε -mate, and the Gap-Tail lemma can be applied to propagate the bound.

\mathcal{U} -family consists of α -mates and β -mates

Illustration of $\alpha+\beta$ -family

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It is more complicated to bound the size of a $\alpha+\beta$ -family:

$$|\alpha+\beta\text{-family}| \leq \begin{cases} \lceil \frac{u(1)-u(2)}{2} \rceil |u_1| & \text{if } u(2) = 1 \\ \frac{u(1)-u(2)}{2} |u_1| & \text{if } u(2) > 1 \end{cases}$$

- Either there are no other double squares, and then it can be shown directly that the bound holds, or
- There is a \mathcal{V} underneath, and we can use induction using the Gap-Tail lemma. \mathcal{V} must be either δ -mate, or ε -mate, and the Gap-Tail lemma can be applied to propagate the bound. (*Special care needed for ε -mate case and super- ε -mate must be put in play !*)

\mathcal{U} -family consists of α -mates, β -mates, and γ -matesIllustration of $\alpha+\beta+\gamma$ -family

R_1		
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aabaabaabaabaabaabaabaabaabaabaaba		5 1 <--- start of α -segment
[][] ()	
abaabaabaabaabaabaabaabaabaabaaba		5 1 <--- end of α -segment
[][] ()	
aabaabaabaabaabaabaabaabaabaabaaba		4 2 <--- start of β -segment
[][] ()	
abaabaabaabaabaabaabaabaabaabaaba		4 2 <--- end of β -segment
[][] ()	
aabaabaabaabaabaabaabaabaabaabaaba		3 3 <--- start of γ -segment
[][] ()	
abaabaabaabaabaabaabaabaabaabaaba		3 3
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baabaabaabaabaabaabaabaabaabaaba		3 3
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baabaabaabaabaabaabaabaabaabaaba		2 4 not a double square
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aabaabaabaabaabaabaabaabaabaabaaba		2 4 not a double square
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abaabaabaabaabaabaabaabaabaabaaba		1 5 not a double square
[][] ()	
baabaabaabaabaabaabaabaabaabaaba		1 5 not a double square

 R_1

It is quite complex to bound the size of a $\alpha+\beta+\gamma$ -family:

$$|\alpha+\beta+\gamma\text{-family}| \leq \frac{2}{3}(u(1) + 1)|u_1|$$

- Either there are no other double squares, and then it can be shown directly that the bound holds, or
- There is a \mathcal{V} underneath, and we can use induction using the Gap-Tail lemma. \mathcal{V} must be either δ -mate, or ε -mate, and the Gap-Tail lemma can be applied to propagate the bound.

Main theorems

Theorem

The number of double squares in a string of length n is bounded by $\lfloor 5n/6 \rfloor$.

Corollary

The number of distinct squares in a string of length n is bounded by $\lfloor 11n/6 \rfloor$.

- We presented a universal upper bound of $\frac{11n}{6}$ for the maximum number of distinct squares in a string of length n
- A bound of $\frac{5n}{6}$ for the maximum number of double squares
- It improves the universal bound of $2n$ by Fraenkel-Simpson
- It improves the asymptotic bound of $2n - \Theta(\log n)$ by Ilie
- The combinatorics of double squares is interesting on its own and possibly can be used for some other problems

THANK YOU



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