# How many double squares can a string contain?

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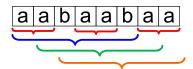


### Motivation and background

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.



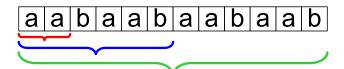
has 6 occurrences of squares, but only 4 distinct squares, *aa*, *aabaab*, *abaaba*, and *baabaa*.



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?

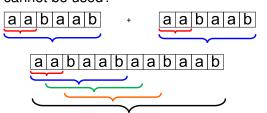
Why this is not simple? In a string of length n,  $O(\log n)$ squares can start at the same position!







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,  $|w| \ge |u| + |v|$ .





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) = 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 denotes 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 > 2.

*primitive root* of x is the smallest primitive y so that  $x = y^n$ .

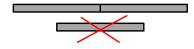
x and y are conjugates if x = uv and y = vu for some u, v.





#### Lemma (Synchronization principle)

Given a primitive string x, a proper suffix y of x, a proper prefix z of x, and  $m \ge 0$ , 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. In particular, if x and y are primitive, then x and y are conjugates.



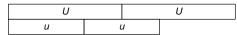
### Double squares

- 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

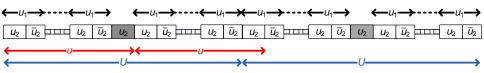
- (b) both uu and UU are rightmost occurrences, or
- (a) |U| < |uu| and either uu or UU is primitively rooted.

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  $1 \le e_2 \le e_1$  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.





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



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



Cyclic shift (rotation) to the right is controlled by

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

while cyclic shift to the left is controlled by

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

lcp = largest common prefix
lcs = largest common suffix



$$u_1 = aaabaa$$
,  $u_2 = aaab$ ,  $\overline{u}_2 = aa$ ,  $u(1) = u(2) = 2$ 

#### 



$$u_1 = aaabaa$$
,  $u_2 = aaab$ ,  $\overline{u}_2 = aa$ ,  $u(1) = 2$ , and  $u(2) = 1$ .

#### 

[	][	) (	]	)			
aaabaaaaabaa	aaabaaab	aa <mark>aaaba</mark>	aaaabaaaaaa	<u>baaab</u> aa			
[	][	) (	]	)			
. <u>aabaaaaaba</u> aa <u>aabaaaba</u> aa <mark>aabaaaaaba</mark> aa <u>aabaaaba</u> aa							
[	][	) (	]	)			
<u>abaaaaabaa</u> aa <u>abaaabaa</u> aa <u>abaaabaa</u> aa <u>abaaabaa</u> aa .							
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baaaaabaaaaabaaabaaaaabaaaabaaaab							



#### Definition

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

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

$$u_1^{\,(\mathcal{U}(1)-1)}u_2\overline{u}_2^{\,}u_2^{\,}u_2^{\,}\overline{u}_2^{\,}u_1^{\,}{}^{\mathcal{U}(2)+\mathcal{U}(1)-2}u_2\overline{u}_2^{\,}u_2^{\,}u_2^{\,}\overline{u}_2^{\,}u_1^{\,}{}^{(\mathcal{U}(2)-1)}$$

N₂ natural inversion factors





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



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  $U: u = u_1^{U(1)}u_2$  and  $U = u_1^{U(1)}u_2u_1^{U(2)}$ .

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

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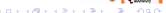
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If the inversion factor in the second v were from  $[L_2, R_2]$ , then |v| = |U|, a contradiction.

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

Therefore vv contains another copy of  $u_1^{\mathcal{U}(1)}u_2u_1^{\mathcal{U}(1)}u_2 = uu$ , 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  $\mathfrak{s}(\mathcal{U}) < \mathfrak{s}(\mathcal{V})$ , then either

- (a)  $\mathfrak{s}(\mathcal{V}) < R_1(\mathcal{U})$ , in which case either
  - (a<sub>1</sub>) V is an  $\alpha$ -mate of U (cyclic shift), or
  - $(a_2)$  V is a  $\beta$ -mate of U (cyclic shift of U to V), or
  - (a<sub>3</sub>) V is a  $\gamma$ -mate of U (cyclic shift of U to v), or
  - (a<sub>4</sub>) V is a  $\delta$ -mate of U (big tail),

or

- (b)  $R_1(\mathcal{U}) \leq \mathfrak{s}(\mathcal{V})$ , then
  - (b<sub>1</sub>) V is a  $\varepsilon$ -mate of U (big gap).





#### $\alpha$ -mate (cyclic shift):

```
R_1
```

```
aaabaaaaabaaabaaabaaaaabaaaabaaabaa
```



#### $\beta$ -mate (cyclic shift of *U* to *V*)



```
baaaaabaaaabaaaaabaaaaabaaa abaaabaaaa baaaaabaaaaabaaaabaaaaabaaaa baaabaaaa
```



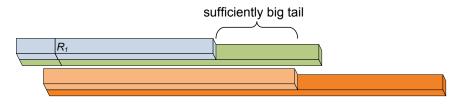
#### $\gamma$ -mate (cyclic shift of *U* to *v*)



```
][ )(
aabaabaabaabaabaabaabaabaabaab
                      ][
                               ) (
```

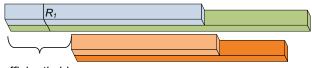


#### $\delta$ -mate (big tail)





#### $\varepsilon$ -mate (big gap)



sufficiently big gap

aabaabaabaabaabaabaabaabaabaabaab 1[



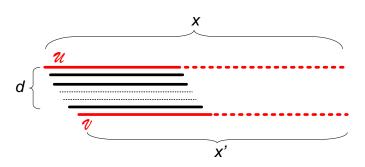
### An upper bound for the number of double squares

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  $G(\mathcal{U}, \mathcal{V})$  is "big" or the tail  $T(\mathcal{U}, \mathcal{V})$  is "big" (for  $\delta$ -mate and  $\varepsilon$ -mate), or it is case of  $\alpha$ -mate,  $\beta$ -mate, or  $\gamma$ -mate.





### Lemma (Gap-Tail lemma)

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

$$\delta(x) \leq \frac{5}{6}|x| - \frac{1}{3}|u| + d - \frac{1}{2}|G(\mathcal{U}, \mathcal{V})| - \frac{1}{3}|T(\mathcal{U}, \mathcal{V})|$$







We deal with  $\alpha$ -mates,  $\beta$ -mates, and  $\gamma$ -mates separately.

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



### $\mathcal{U}$ -family consists only of $\alpha$ -mates

Illustration of  $\alpha$ -family with  $\mathcal{U}(1) = \mathcal{U}(2)$ 

baaaaabaaa abaaabaaa abaaaabaaaaabaaa abaaabaaa



#### Illustration of $\alpha$ -family with $\mathcal{U}(1) > \mathcal{U}(2)$

#### 

[	][	) (	]	)
aaabaaaaabaa	aaabaaab	aaaaba	aaaabaa <u>aaa</u>	<u>baaab</u> aa
[	][	) (	]	)
. aabaaaaaba	aabaaab	aaaaaba	aaaabaaaa <u>aa</u>	<u>baaaba</u> aa
[	][	) (	]	)
abaaaaabaa	aa <mark>abaa</mark> ab	aaaaaba	aaaabaa aa <u>a</u>	<u>baa</u> abaaaa.
[	][	) (	]	)
baaaaabaa	aaabaaab	aaaaaba	aaaabaaaaa	baaabaaa



It is easy to estimate the size of  $\alpha$ -family, as it is controlled by  $lcp(u_1, \overline{u}_1)$  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  $\gamma$ -mate, or  $\delta$ -mate, or  $\varepsilon$ -mate, and the Gap-Tail lemma can be applied to propagate the bound.



### $\mathcal{U}$ -family consists of $\alpha$ -mates and $\beta$ -mates

#### Illustration of $\alpha+\beta$ -family

```
aaabaaaaabaaaabaaaabaaaaab aaaabaaab aaabaaabaaaabaaaabaaaabaaaabaaaab aaabaaab aaabaaaabaaaaabaaaaa
aaabaaaaabaaaaabaaaaabaaaaab aaabbaaab aaabaaaaabaaaaabaaaaabaaaaabaaaaab aaabaaaab
aaabaaaaabaaaaabaaaaab aaabaaab aaabaaaaabaaaaabaaaaab aaabaaab aaabaa...
     ...abaaaaabaaaaabaaaaabaa ahaaabaa abaaaaabaaaaabaaaaabaa ahaaaaabaa abaaaa
    . .baaaaabaaaabaaaabaaa baaabaaa baaaaabaaaaabaaaaabaaaabaaa baaabaaa baaaaa
       ...abaaaaabaaaaabaa abaaabaa abaaaaabaaaaabaaaaabaaaabaaa abaaabaa
       ....baaaaabaaaaabaaa baaabaaa baaaaabaaaaabaaaaabaaaaabaaa baaabaaa baaaabaaaa
```

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

$$\leq \begin{cases} \left\lceil \frac{\mathcal{U}(1) - \mathcal{U}(2)}{2} \right\rceil |u_1| & \text{if } \mathcal{U}(2) = 1 \\ \\ \frac{\mathcal{U}(1) - \mathcal{U}(2)}{2} |u_1| & \text{if } \mathcal{U}(2) > 1 \end{cases}$$

- Either there are no other double squares, and then it can be shown directly that the bound holds, or
- There is a V underneath, and we can use induction using the Gap-Tail lemma. 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 $\alpha$ -mates, $\beta$ -mates, and $\gamma$ -mates

#### Illustration of $\alpha+\beta+\gamma$ -family

```
aabaabaabaabaabaabaabaabaabaabaabaab
                      <--- start of a-segment
abaabaabaabaabaabaabaabaabaabaaba
 aabaabaabaabaabaabaabaabaabaabaabaab
                       <--- start of 8-segment
 abaabaabaabaabaabaabaabaabaabaabaaba
                       <--- end of 8-segment
  2 4 not a double square
    2 4 not a double square
                                 not a double square
     not a double square
     1 5 not a double square
     not a double square
```





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

$$\leq \frac{2}{3} \big( \, \mathcal{U}(1) + 1 \big) | \mathcal{U}_1 |$$

- 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  $\delta$ -mate, or  $\varepsilon$ -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 |5n/6|.

#### Corollary

The number of distinct squares in a string of length n is bounded by |11n/6|.





- We presented a universal upper bound of <sup>11n</sup>/<sub>6</sub> 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  $2n \Theta(n)$  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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