

#### **CIBB 2019**



# Modification of Valiant's Parsing Algorithm for String-Searching Problem

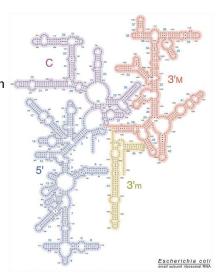
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JetBrains Research, Programming Languages and Tools Lab Saint Petersburg University

September 6, 2019

#### RNA Analysis

- RNA secondary structure prediction
- Applications: classification and recognition problems
- String-searching problem

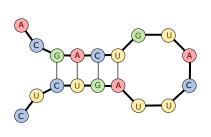


#### Formal Grammars and Languages

- $G = (\Sigma, N, R, s)$  context-free grammar (CFG) in Chomsky normal form
  - ightharpoonup a 
    ightharpoonup bc, where  $a, b, c \in N$
  - ▶  $a \rightarrow A$ , where  $a \in N, A \in \Sigma$
  - $s \to \varepsilon$ , where  $\varepsilon$  is an empty string
- $L_G(s) = \{\omega \mid s \Rightarrow^* \omega\}$ , where  $\omega \in \Sigma^*$
- Parsing does  $\omega$  belong to  $L_G(s)$ ?

#### CFG-based Approach

- RNA sequences are treated as strings over  $\Sigma = \{A, G, C, U\}$ CACGACUGUACUUAGUCUC...CUGGAUCACCUCCUU
- CFG describes RNA secondary structure features

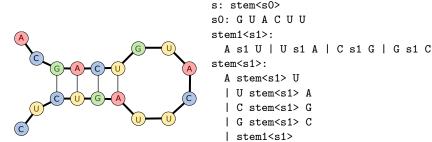


```
s: stem<s0>
s0: G U A C U U
stem1<s1>:
    A s1 U | U s1 A | C s1 G | G s1 C
stem<s1>:
    A stem<s1> U
    | U stem<s1> A
    | C stem<s1> C
```

stem1 < s1 >

#### CFG-based Approach

- RNA sequences are treated as strings over  $\Sigma = \{A, G, C, U\}$ CACGACUGUACUUAGUCUC...CUGGAUCACCUCCUU
- CFG describes RNA secondary structure features



- Parsing as method to find all substrings with specific secondary structure features
- String-searching problem: for input string of length  $n = 2^p 1$  find all substrings of length m which belong to  $L_G(s)$

#### **Problems**

- Long sequences
   Large amount of data
   Complex models
- $\implies$  development of efficient parsing algorithms

#### Tabular Parsing Algorithms

- Input:
  - Grammar  $G = (\Sigma, N, R, s)$  in Chomsky normal form
  - ▶ String  $\omega = \omega_1 \omega_2 \dots \omega_n$ ,  $\omega_i \in \Sigma$
- Parsing table T:
  - $T_{i,j} = \{a \mid a \in \mathbb{N}, \omega_{i+1} \dots \omega_i \in L_G(a)\} \quad \forall i < j\}$
  - $\omega \in L_G(s) \iff s \in T_{0,n}$

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- Process of filling:
  - $T_{i-1,i} = \{ a \mid a \to \omega_i \in R \}$
  - ▶  $T_{i,j} = f(P_{i,j})$ , where  $P_{i,j} = \bigcup_{k=i+1}^{j-1} T_{i,k} \times T_{k,j}$  $f(P_{i,j}) = \{a \mid \exists a \rightarrow bc \in R : (b,c) \in P_{i,j}\}$

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#### To Valiant's Parsing Algorithm

CYK: 
$$\mathcal{O}(|G|n^3)$$

Younger, D. H. "Context-free language processing in time  $n^3$ " 1966

 $\Downarrow$ 

Reduction to matrix multiplication



Reduction to Boolean matrix multiplication



Valiant:  $\mathcal{O}(|G|BMM(n)log(n))$ 

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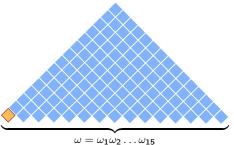
Reduction to Boolean matrix multiplication



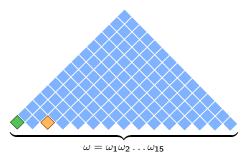
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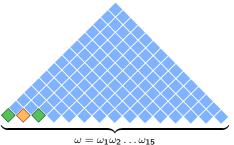
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  - ▶ Utilization of parallel techniques and highly-efficient libraries
  - Generalization to more powerful classes of formal grammars: conjunctive and Boolean
- **●** (−):
  - ▶ Not suitable for string-searching problem It is necessary to calculate at least 2 triangle submatrices of size  $\frac{n}{2}$



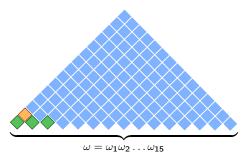
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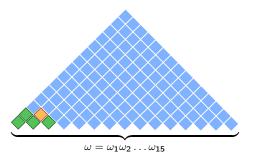
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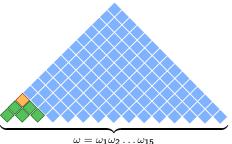
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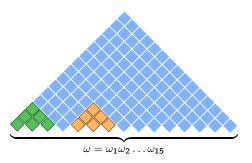


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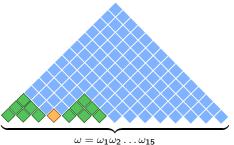


Parsing for String-Searching Problem

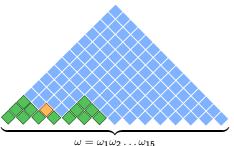
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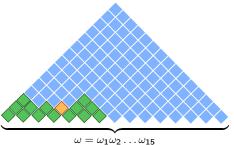
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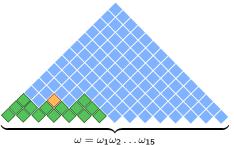
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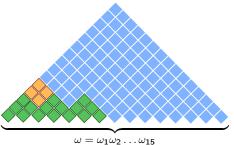
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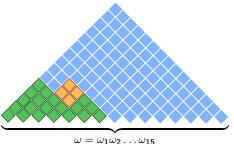
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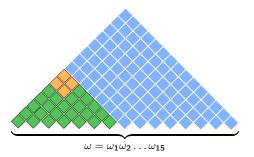
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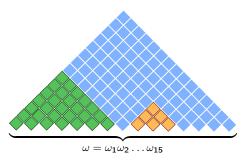
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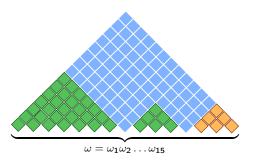
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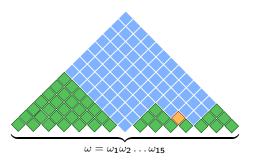
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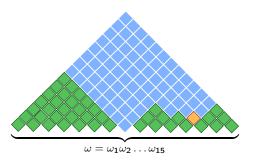
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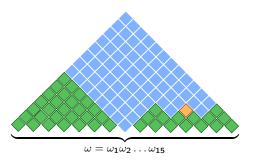
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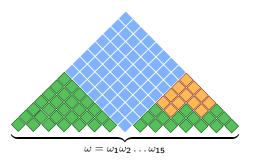
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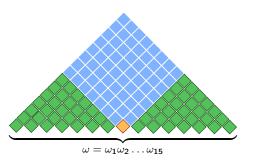
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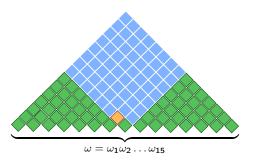
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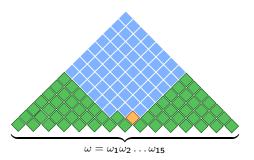
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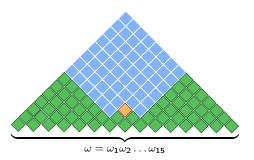
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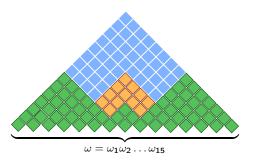
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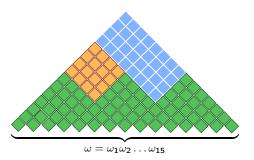
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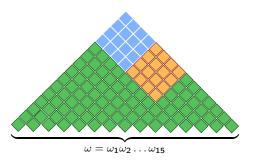
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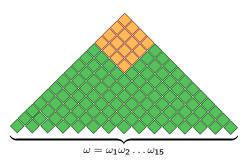
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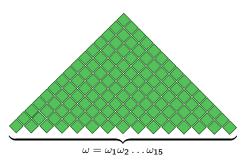
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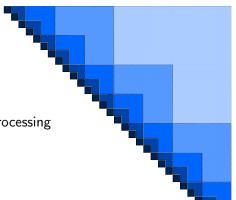


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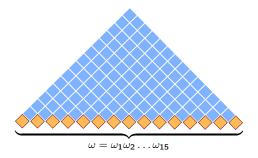
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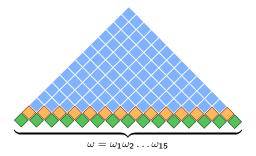


- Rearranging the submatrices processing
- Division the parsing table into layers of disjoint submatrices

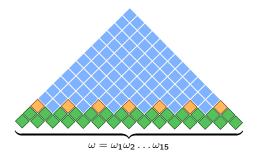
- Each matrix in the layer can be handled independently
- Increasing the lever of parallelism:
  - Matrix multiplication
  - ► Each matrix in layer
  - ► Each pair of nonterminals



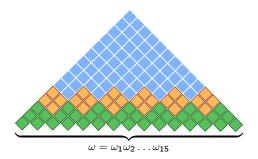
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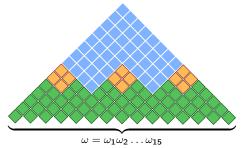
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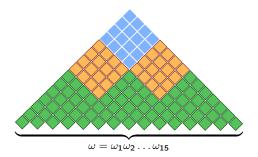
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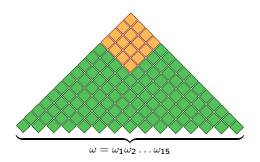
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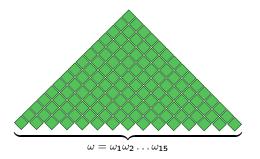
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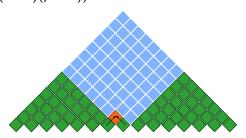


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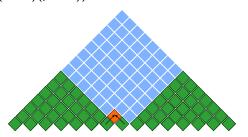
#### Application for the String-Searching Problem

- **Problem**: for input string of length  $n = 2^p 1$  find all substrings of length m which belong to  $L_G(s)$
- Valiant's algorithm: it is necessary to calculate at least 2 triangle submatrices of size  $\frac{n}{2}$   $\mathcal{O}(|G|BMM(2^{p-1})(p-2))$



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• Modification: it is necessary to compute layers with submatrices of size not greater than  $2^r$ , where  $2^{r-2} < s \le 2^{r-1}$   $\mathcal{O}(|G|2^{2(p-r)-1}BMM(2^r)(r-1))$ 

#### **Evaluation**

Implementation

► CPU-based: M4RI library

GPU-based: CUDA C

Grammars

```
► D<sub>2</sub>:
                     s: ss | (s) | [s] | \varepsilon
▶ BIO:
                     s: stem<s0>
                     any_str: any_smb*[2..10]
                     s0: any_str | any_str stem<s0> s0
                     any_smb: A | T | C | G
                     stem1<s1>: A s1 T | G s1 C | T s1 A | C s1 G
                     stem2<s1>: stem1<stem1<s1>>
                     stem<s1>:
                           A stem<s1> T
                         | T stem<s1> A
                           C stem<s1> G
                         | G stem<s1> C
                           stem1<stem2<s1>>
```

#### Data preprocessing

• Example: n = 15, m = 6

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AAGCTTGAAGCTTG len = 15

## Results: Comparative Analysis

	Time (sec)								
n	Grammar $D_2$			Grammar <i>BIO</i>					
	valCPU	modCPU	valGPU	modGPU	valCPU	modCPU	valGPU	modGPU	
127	0.08	0.08	0.20	0.10	1.35	1.34	0.19	0.10	
255	0.28	0.30	0.52	0.13	5.40	5.50	0.53	0.14	
511	1.21	1.18	1.90	0.25	21.97	22.35	1.99	0.26	
1023	4.90	4.78	7.88	0.54	88.70	90.32	7.89	0.60	
2047	19.61	19.38	33.50	1.50	363.32	374.20	34.01	1.70	
4095	78.36	78.28	140.47	4.45	1467.68	1480.59	141.10	5.47	
8191	315.67	315.08	-	13.65	-	-	-	18.04	

## Results: Comparative Analysis

		Time (sec)							
n	Grammar D <sub>2</sub>			Grammar <i>BIO</i>					
	valCPU	modCPU	valGPU	modGPU	valCPU	modCPU	valGPU	modGPU	
127	0.08	0.08	0.20	0.10	1.35	1.34	0.19	0.10	
255	0.28	0.30	0.52	0.13	5.40	5.50	0.53	0.14	
511	1.21	1.18	1.90	0.25	21.97	22.35	1.99	0.26	
1023	4.90	4.78	7.88	0.54	88.70	90.32	7.89	0.60	
2047	19.61	19.38	33.50	1.50	363.32	374.20	34.01	1.70	
4095	78.36	78.28	140.47	4.45	1467.68	1480.59	141.10	5.47	
8191	315.67	315.08	-	13.65	-	-	-	18.04	

## Results: Comparative Analysis

	Time (sec)								
n	Grammar $D_2$			Grammar <i>BIO</i>					
	valCPU	modCPU	valGPU	modGPU	valCPU	modCPU	valGPU	modGPU	
127	0.08	0.08	0.20	0.10	1.35	1.34	0.19	0.10	
255	0.28	0.30	0.52	0.13	5.40	5.50	0.53	0.14	
511	1.21	1.18	1.90	0.25	21.97	22.35	1.99	0.26	
1023	4.90	4.78	7.88	0.54	88.70	90.32	7.89	0.60	
2047	19.61	19.38	33.50	1.50	363.32	374.20	34.01	1.70	
4095	78.36	78.28	140.47	4.45	1467.68	1480.59	141.10	5.47	
8191	315.67	315.08	-	13.65	-	-	-	18.04	

## Results: String-searching Problem

m		Time (sec)						
m	n	valCPU	modCPU	valGPU	modGPU			
250	1023	4.90	3.00	7.88	0.24			
	2047	19.61	6.65	33.50	0.26			
250	4095	78.36	13.83	140.47	0.32			
	8191	315.67	28.90	-	0.46			
	2047	19.61	12.18	33.50	0.58			
510	4095	78.36	26.58	140.47	0.65			
	8191	315.67	56.70	-	0.88			
1020	4095	78.36	48.31	140.47	1.59			
	8191	315.67	108.38	_	1.95			
2040	8191	315.67	197.32	-	5.10			

#### Conclusion

- The modification of Valiant's algorithm was proposed
  - Layered submatrices processing
  - Effective utilization of parallel techniques and GPGPU
- The modification is applicable to the string-searching problem

#### Future Research

- Improvement of the existing implementation
- Evaluation on real-world data
- Extension for more expressive classes of formal languages (conjunctive, Boolean)

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Thanks!