

Analytic Pattern Matching: From DNA to Twitter

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Combinatorial Pattern Matching, Ischia Island, 2015

*Joint work with Philippe Jacquet

Outline

1. Motivations

- Finding Biological Signals
- Searching Google
- Classifying Twitter

2. Pattern Matching Problems

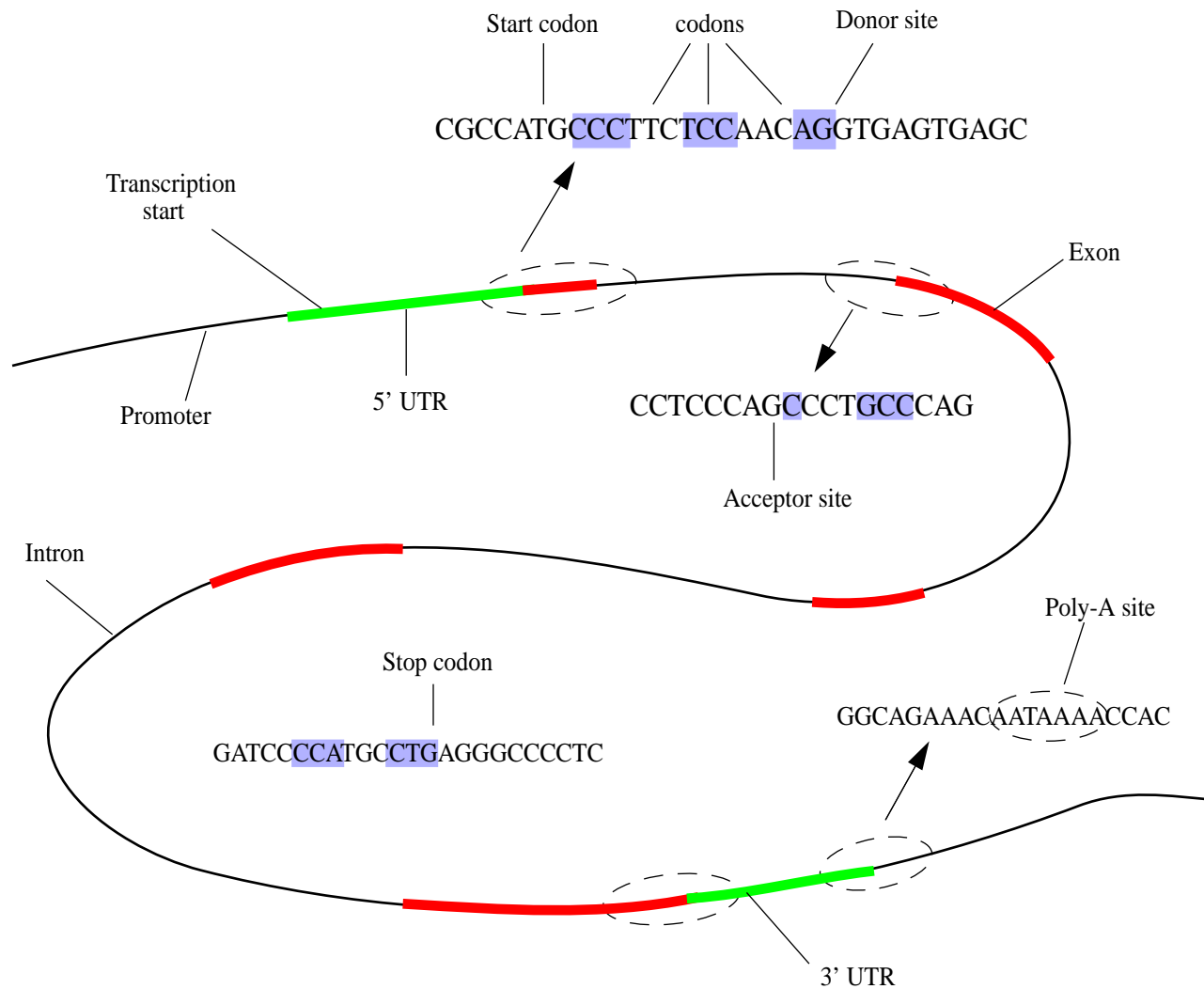
- Exact String Matching
- Constrained String Matching
- Generalized String Matching
- Subsequence String Matching
- String Complexity

3. Analysis & Applications

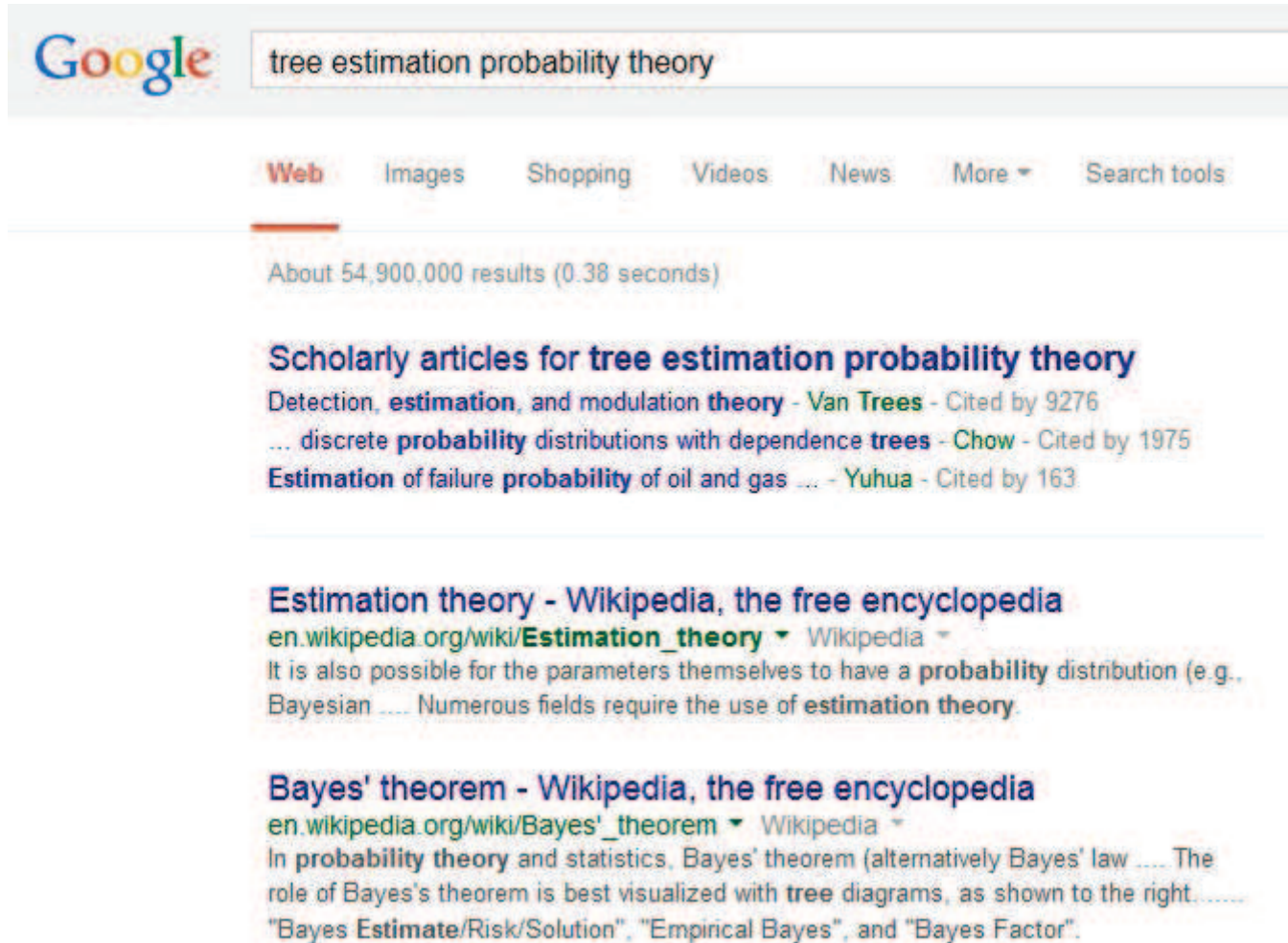
- Exact String Matching & Finding Biological Motifs
- Hidden Patterns & Intrusion Detection
- Joint String Complexity & Classification of Twitter

Motivation – Biology & String Matching

Biological world is highly **stochastic** and **inhomogeneous** (S. Salzberg).



Motivation – Google & Subsequence Matching



The image is a screenshot of a Google search results page. At the top, the Google logo is on the left, and the search query 'tree estimation probability theory' is in the search bar. Below the search bar, there are tabs for 'Web', 'Images', 'Shopping', 'Videos', 'News', 'More', and 'Search tools'. The 'Web' tab is selected. Below the tabs, it says 'About 54,900,000 results (0.38 seconds)'. The first result is titled 'Scholarly articles for tree estimation probability theory' and lists three articles: 'Detection, estimation, and modulation theory - Van Trees - Cited by 9276', '... discrete probability distributions with dependence trees - Chow - Cited by 1975', and 'Estimation of failure probability of oil and gas ... - Yuhua - Cited by 163'. The second result is titled 'Estimation theory - Wikipedia, the free encyclopedia' and includes the URL 'en.wikipedia.org/wiki/Estimation_theory' and a snippet: 'It is also possible for the parameters themselves to have a probability distribution (e.g., Bayesian Numerous fields require the use of estimation theory.' The third result is titled 'Bayes' theorem - Wikipedia, the free encyclopedia' and includes the URL 'en.wikipedia.org/wiki/Bayes'_theorem' and a snippet: 'In probability theory and statistics, Bayes' theorem (alternatively Bayes' law The role of Bayes's theorem is best visualized with tree diagrams, as shown to the right..... "Bayes Estimate/Risk/Solution", "Empirical Bayes", and "Bayes Factor".'

Google

tree estimation probability theory

Web Images Shopping Videos News More Search tools

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Motivation – Intrusion Detection & Hidden Patterns

Convert all color commands to black or white. Since PostScript files are often extremely large, it makes sense to try to compress them with either the zip or gzip programs. In such a case, the eps file is replaced by a file with extension zip or eps.gz, or eps.gz. Two problems now arise: first LATEX cannot read such files to obtain the bounding box information, and secondly, the driver needs to unpack such a file to include it in the final output. This can be accomplished with, for example: `DeclareGraphicsRule.eps.gzeps.eps.bbgunzip` which stabilizes the graphics type as eps with the bounding box information in the file of the same name and extension. Convert all color commands to black or white.

How to know whether this subsequence observed in an audit file constitutes an attack or is merely a result of randomness?

Goal: Minimize the number of false positives!

Motivation – Twitter & String Complexity

"allow users to download an entire movie in one second." I need this <http://t.co/3fbNfKEkah>

Green energy boss accuses Govt of obstructing renewable energy development <http://t.co/v5Lq2Jx1GQ>

Figure 1: Two similar twitter texts have many common words

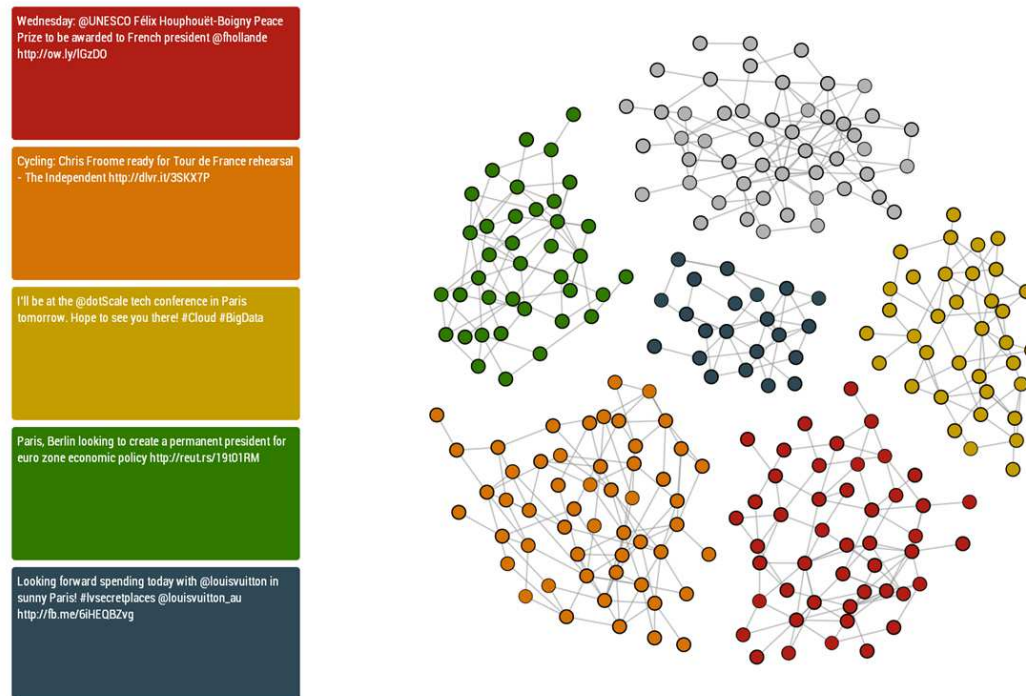


Figure 2: Twitters Classification

Outline Update

1. Motivations
2. Pattern Matching Problems
 - Exact String Matching
 - Constrained String Matching
 - Generalized String Matching
 - Subsequence String Matching
 - String Complexity
3. Analysis & Applications

Pattern Matching

Let \mathcal{W} and T be (set of) strings generated over a finite alphabet \mathcal{A} .

We call \mathcal{W} the **pattern** and T the **text**. The text T is of length n and is generated by a **probabilistic source**.

Text: $T_m^n = T_m \dots T_n$.

Pattern: $\mathcal{W} = w_1 \dots w_m, w_i \in \mathcal{A}$;

Set of patterns: $\mathcal{W} = \{\mathcal{W}_1, \dots, \mathcal{W}_d\}$ with $\mathcal{W}_i \in \mathcal{A}^{m_i}$.

Basic question:

how many times \mathcal{W} occurs in T (or how long to wait until \mathcal{W} occurs in T).

Define

$$O_n(\mathcal{W}) = \#\{i : T_{i-m+1}^i = \mathcal{W}, m \leq i \leq n\}$$

as the number of w occurrences in the text T_1^n .

Our goal: Study **probabilistic behavior** of $O_n(\mathcal{W})$ for various pattern matching problems using tools of **analytic combinatorics**.

Variations on Pattern Matching

(Exact) String Matching: In the exact string matching the pattern

$$\mathcal{W} = w_1 \dots w_m$$

is a **given string** (i.e., consecutive sequence of symbols).

Generalized String Matching: In the generalized pattern matching a **set of patterns** (rather than a single pattern) is given, that is,

$$\mathcal{W} = (\mathcal{W}_0, \mathcal{W}_1, \dots, \mathcal{W}_d), \quad \mathcal{W}_i \in \mathcal{A}^{m_i}$$

where \mathcal{W}_i itself for $i \geq 1$ is a subset of \mathcal{A}^{m_i} .

The set \mathcal{W}_0 is called the **forbidden set**.

Three cases to be considered:

$\mathcal{W}_0 = \emptyset$ — interest in the number of patterns from \mathcal{W} occurring in the text.

$\mathcal{W}_0 \neq \emptyset$ — we study the number of $\mathcal{W}_i, i \geq 1$ pattern occurrences **under the condition** that no pattern from \mathcal{W}_0 occurs in the text.

$\mathcal{W}_i = \emptyset, i \geq 1, \mathcal{W}_0 \neq \emptyset$ — restricted pattern matching.

Pattern Matching Problems

Hidden Words or Subsequence Pattern Matching: We search for a subsequence $\mathcal{W} = w_1 \dots w_m$ rather than a string in a text.

That is, there are indices $1 \leq i_1 < i_2 < \dots < i_m \leq n$ such that

$$T_{i_1} = w_1, T_{i_2} = w_2, \dots, T_{i_m} = w_m.$$

We also say that the word \mathcal{W} is “hidden” in the text.

For example:

$\mathcal{W} = \text{date}$

$T = \text{hidden pattern}$

occurs four times as a subsequence in the text as hidden pattern.

Joint String Complexity: For a given string X , we ask how many distinct subwords it contains. This is called string complexity.

For two strings X and Y , we want to know how many common and distinct subwords they contain. This is called joint string complexity.

New Book on Pattern Matching

How do you distinguish a cat from a dog by their DNA?
Did Shakespeare really write all of his plays?

Pattern matching techniques can offer answers to these questions and to many others, from molecular biology, to telecommunications, to classifying Twitter content.

This book for researchers and graduate students demonstrates the probabilistic approach to pattern matching, which predicts the performance of pattern matching algorithms with very high precision using analytic combinatorics and analytic information theory. Part I compiles known results of pattern matching problems via analytic methods. Part II focuses on applications to various data structures on words, such as digital trees, suffix trees, string complexity and string-based data compression. The authors use results and techniques from Part I and also introduce new methodology such as the Mellin transform and analytic depoissonization.

More than 100 end-of-chapter problems help the reader to make the link between theory and practice.

Philippe Jacquet is a research director at INRIA, a major public research lab in Computer Science in France. He has been a major contributor to the Internet OLSR protocol for mobile networks. His research interests involve information theory, probability theory, quantum telecommunication, protocol design, performance evaluation and optimization, and the analysis of algorithms. Since 2012 he has been with Alcatel-Lucent Bell Labs as head of the department of Mathematics of Dynamic Networks and Information. Jacquet is a member of the prestigious French Corps des Mines, known for excellence in French industry, with the rank of "Ingenieur General". He is also a member of ACM and IEEE.

Wojciech Szpankowski is Saul Rosen Professor of Computer Science and (by courtesy) Electrical and Computer Engineering at Purdue University, where he teaches and conducts research in analysis of algorithms, information theory, bioinformatics, analytic combinatorics, random structures, and stability problems of distributed systems. In 2008 he launched the interdisciplinary Institute for Science of Information, and in 2010 he became the Director of the newly established NSF Science and Technology Center for Science of Information. Szpankowski is a Fellow of IEEE and an Erskine Fellow. He received the Humboldt Research Award in 2010.

Cover design: Andrew Ward

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Jacquet and
Szpankowski

Analytic Pattern Matching

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Analytic Pattern Matching

From DNA to Twitter

#STRINGS

#ASYMPTOT

#PROBA

#COMBINATOR

#TEXTS

COMPLEXITY

MARKOV

ATGCATTAGCTACGT

ATGCATTAGCTACGT

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Book Contents

Chapter 1: Probabilistic Models

Chapter 2: Exact String Matching

Chapter 3: Constrained Exact String Matching

Chapter 4: Generalized String Matching

Chapter 5: Subsequence String Matching

Chapter 6: Algorithms and Data Structures

Chapter 7: Digital Trees

Chapter 8: Suffix Trees & Lempel-Ziv'77

Chapter 9: Lempel-Ziv'78 Compression Algorithm

Chapter 10: String Complexity

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Analysis: Exact String Matching

Memoryless Source: The text is generated by a memoryless source with probability of seeing $a \in \mathcal{A}$ equal to $P(a)$.

Tools. Symbolic calculus and analytic tools of languages:

- (i) Language \mathcal{L} is a collection of words satisfying some properties.
- (ii) Generating function $L(z)$ of language \mathcal{L} is defined as

$$L(z) = \sum_{u \in \mathcal{L}} P(u) z^{|u|}.$$

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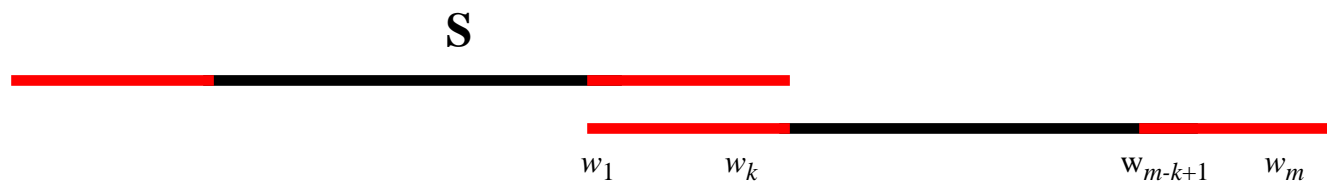
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Autocorrelation Polynomial: For \mathcal{W} define the autocorrelation set \mathcal{S} as:

$$\mathcal{S} = \{w_{k+1}^m : w_1^k = w_{m-k+1}^m\},$$

and \mathcal{WW} is the set of positions k satisfying $w_1^k = w_{m-k+1}^m$.



The generating function $S(z)$ of \mathcal{S} is called the autocorrelation polynomial:

$$S(z) = \sum_{k \in \mathcal{WW}} P(w_{k+1}^m) z^{m-k}.$$

Example: For $\mathcal{W} = bab$, we have $\mathcal{WW} = \{1, 3\}$ and $\mathcal{S} = \{\epsilon, ab\}$.

Language \mathcal{T}_r and Associated Languages

Define \mathcal{T}_r as set of words containing exactly $r \geq 1$ occurrences of \mathcal{W} :

$$\mathcal{T}_r = \mathcal{R} \cdot \mathcal{M}^{r-1} \cdot \mathcal{U}.$$

which can be illustrated for \mathcal{T}_4 as follows



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- (i) Language \mathcal{R} : set of words containing only one occurrence of \mathcal{W} , located at the right end. **For example:** for $\mathcal{W} = aba$, we have $ccaba \in \mathcal{R}$.
- (ii) Language \mathcal{U} :

$$\mathcal{U} = \{u : \mathcal{W} \cdot u \in \mathcal{T}_1\}$$

that is, a word $u \in \mathcal{U}$ if $\mathcal{W} \cdot u$ has exactly one occurrence of \mathcal{W} at the left end of $\mathcal{W} \cdot u$. **For example:** $bba \in \mathcal{U}$ (since $ababba \in \mathcal{T}_1$) but $ba \notin \mathcal{U}$

- (iii) Language \mathcal{M} :

$$\mathcal{M} = \{u : \mathcal{W} \cdot u \in \mathcal{T}_2 \text{ and } \mathcal{W} \text{ occurs at the right of } \mathcal{W} \cdot u\},$$

that is, \mathcal{M} is a language such that $\mathcal{W}\mathcal{M}$ has exactly two occurrences of \mathcal{W} at the left and right end of a word from \mathcal{M} .

For example: $ba \in \mathcal{M}$ since $ababa$).

Language Relations & Generating Functions

Lemma 1. (i) *The languages \mathcal{M}, \mathcal{U} and \mathcal{R} satisfy:*

$$\begin{aligned}\mathcal{U} \cdot \mathcal{A} &= \mathcal{M} + \mathcal{U} - \{\epsilon\}, \\ \bigcup_{k \geq 1} \mathcal{M}^k &= \mathcal{A}^* \cdot \mathcal{W} + \mathcal{S} - \{\epsilon\}, \quad \mathcal{W} \cdot \mathcal{M} = \mathcal{A} \cdot \mathcal{R} - (\mathcal{R} - \mathcal{W}),\end{aligned}$$

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(ii) *The generating functions associated with languages \mathcal{M}, \mathcal{U} and \mathcal{R} satisfy*

$$\begin{aligned}U_{\mathcal{W}}(z) &= \frac{M(z) - 1}{z - 1} \\ \frac{1}{1 - M(z)} &= S_{\mathcal{W}}(z) + P(\mathcal{W}) \frac{z^m}{1 - z}, \quad R(z) = P(\mathcal{W}) z^m \cdot U_{\mathcal{W}}(z)\end{aligned}$$

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(iii) The generating functions $T_r(z) = \sum_{n \geq 0} \Pr\{O_n(\mathcal{W}) = r\} z^n$ and $T(z, u) = \sum_{r=1}^{\infty} T_r(z) u^r$ satisfy

$$\begin{aligned}T_r(z) &= R(z) M_{\mathcal{W}}^{r-1}(z) U_{\mathcal{W}}(z), \quad r \geq 1 \\ T(z, u) &= R(z) \frac{u}{1 - uM(z)} U_{\mathcal{W}}(z).\end{aligned}$$

Main Results: Asymptotics

Theorem 1 (Regnier and W.S.). (i) *Moments*. for $n \geq m$ we have

$$\mathbf{E}[O_n(\mathcal{W})] = P(\mathcal{W})(n - m + 1), \quad \mathbf{Var}[O_n(\mathcal{W})] = nc_1 + c_2$$

with

$$c_1 = P(\mathcal{W})(2S(1) - 1 - (2m - 1)P(\mathcal{W})) - (m - 1)(2S(1) - 1) - 2S'(1).$$

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(ii) *Probability*. For $r = O(1)$

$$\Pr\{O_n(\mathcal{W}) = r\} \sim \sum_{j=1}^{r+1} (-1)^j a_j \binom{n}{j-1} \rho_{\mathcal{W}}^{-(n+j)}$$

where $\rho_{\mathcal{W}}$ is the smallest root of $D_{\mathcal{W}}(z) = (1 - z)S_{\mathcal{W}}(z) + z^m P(\mathcal{W}) = 0$.

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Central Limit Law: For $r = \mathbf{E}[O_n] + x\sqrt{\mathbf{Var}O_n}$ for $x = O(1)$

$$\Pr\{O_n(\mathcal{W}) = r\} = \frac{1}{\sqrt{2\pi c_1 n}} e^{-\frac{1}{2}x^2} \left(1 + O\left(\frac{1}{\sqrt{n}}\right) \right).$$

Large Deviations: For Case $r = (1 + \delta)EO_n$ with $a = (1 + \delta)P(\mathcal{W})$ we have

$$\Pr\{O_n(\mathcal{W}) \sim (1 + \delta)EO_n\} = \frac{e^{-(n-m+1)I(a)+\delta a}}{\sigma_a \sqrt{2\pi(n-m+1)}}$$

where $I(a) = a\omega_a + \rho(\omega_a)$ and $\rho(t)$ to be the root of $1 - e^t M_{\mathcal{W}}(e^\rho) = 0$.

Biology – Weak Signals and Artifacts

Denise and Regnier (2002) observed that in biological sequence whenever a word is overrepresented, then its subwords are also overrepresented.

For example, if $\mathcal{W}_1 = AATAAA$, then

$$\mathcal{W}_2 = ATAAAN$$

is also overrepresented.

Overrepresented subwords are called artifact, and it is important to disregard automatically noise created by artifacts.

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This harder question needs a new approach thru Generalized Pattern Matching to show that

$$\mathbf{E}[O_n(\mathcal{W}_2) | O_n(\mathcal{W}_1) = k] \sim \alpha n.$$

When \mathcal{W}_1 is overrepresented α differs significantly from $\mathbf{E}[O_n(\mathcal{W}_2)]$.

Polyadenylation Signals in Human Genes

Beaudoing et al. (2000) studied several variants of the well known AAUAAA polyadenylation signal in mRNA of humans genes.

Using our approach Denise and Regnier (2002) discovered/eliminated all artifacts and found new signals in a much simpler and reliable way.

Hexamer	Obs.	Rk	Exp.	Z-sc.	Rk	Cd.Exp.	Cd.Z-sc.	Rk
AAUAAA	3456	1	363.16	167.03	1			1
AAAUAA	1721	2	363.16	71.25	2	1678.53	1.04	1300
AUAAAA	1530	3	363.16	61.23	3	1311.03	6.05	404
UUUUUU	1105	4	416.36	33.75	8	373.30	37.87	2
AUAAAU	1043	5	373.23	34.67	6	1529.15	12.43	4078
AAAAUA	1019	6	363.16	34.41	7	848.76	5.84	420
UAAAAU	1017	7	373.23	33.32	9	780.18	8.48	211
AUUAAA	1013	1	373.23	33.12	10	385.85	31.93	3
AUAAAG	972	9	184.27	58.03	4	593.90	15.51	34
UAUAUA	922	10	373.23	28.41	13	1233.24	-8.86	4034
UAAAAA	922	11	363.16	29.32	12	922.67	9.79	155
UUAAAA	863	12	373.23	25.35	15	374.81	25.21	4
CAAUAA	847	13	185.59	48.55	5	613.24	9.44	167
AAAAAA	841	14	353.37	25.94	14	496.38	15.47	36
UAAUAU	805	15	373.23	22.35	21	1143.73	-10.02	4068

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Hidden Patterns

In the subsequence pattern or a hidden word \mathcal{W} occurs as a subsequence:

$$T_{i_1} = w_1, T_{i_2} = w_2, \dots, T_{i_m} = w_m.$$

where we put additional constraints that

$$i_{j+1} - i_j \leq d_j.$$

The $I = (i_1, \dots, i_m)$ -tuple is called a position and $\mathcal{D} = (d_1, \dots, d_m)$ constitutes the constraints.

If all d_j are finite, then we have the **constrained problem**.

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If all d_j are finite, then we have the constrained problem.

Let $O_n(\mathcal{W})$ be the number of \mathcal{W} occurrences in T . Observe that

$$O_n(\mathcal{W}) = \sum_I X_I$$

where

$$X_I := 1 \text{ if } \mathcal{W} \text{ occurs at position } I \text{ in } T_n$$

and 0 otherwise.

How to Analyze It – De Bruijn Automata

1. The $(\mathcal{W}, \mathcal{D})$ **constrained subsequence problem** is viewed as the **generalized string matching**.

Example: If $(\mathcal{W}, \mathcal{D}) = a\#_2b$, then $\mathcal{W} = \{ab, aab, abb\}$.

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2. **de Bruijn Automaton**. Let $M = \max\{\text{length}(\mathcal{W})\} - 1$. Define a de Bruijn automaton over \mathcal{B} where

$$\mathcal{B} = \mathcal{A}^M.$$

De Bruijn automaton is built over \mathcal{B} .

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3. Let $b \in \mathcal{B}$ and $a \in \mathcal{A}$. Then the transition from the state b upon scanning symbol a of the text is to $\hat{b} \in \mathcal{B}$ such that

$$ba \mapsto \hat{b} = b_2b_3 \cdots b_Ma.$$

For example

$$\underbrace{abb}_{\mathcal{B}} \underbrace{a}_{\mathcal{A}} \mapsto \underbrace{bba}_{\mathcal{B}}.$$

4. **The Transition Matrix:** $\mathbf{T}(u)$ is a complex-valued transition matrix defined as:

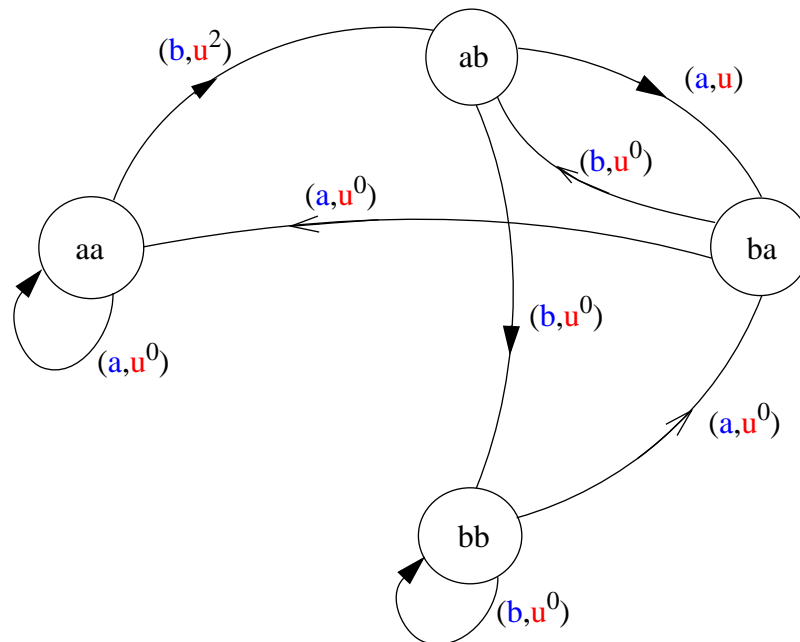
$$[\mathbf{T}(u)]_{b, \hat{b}} := P(a) u^{O_{M+1}(ba) - O_M(b)} \mathbb{I}[\hat{b} = b_2b_3 \cdots b_Ma]$$

where $O_M(b)$ is the number of pattern occurrences \mathcal{W} in the text b .

Example

5. Example. Let $\mathcal{W} = \{ab, aab, aba\}$. Then $M = 2$, and the the de Bruijn graph and matrix $\mathbf{T}(u)$ are shown below

$$\mathbf{T}(u) = \begin{array}{c} \begin{array}{cccc} & aa & ab & ba & bb \end{array} \\ \begin{array}{c} aa \\ ab \\ ba \\ bb \end{array} \left(\begin{array}{ccccc} P(a) & P(b) & u^2 & 0 & 0 \\ 0 & 0 & P(a) & u & P(b) \\ P(a) & P(b) & 0 & 0 & 0 \\ 0 & 0 & P(a) & P(b) & 0 \end{array} \right) . \end{array}$$



Generating Functions

6. Using properties of product of matrices we conclude that

$$O_n(u) = \mathbf{E}[u^{O_n(\mathcal{W})}] = \mathbf{b}^t(u) \mathbf{T}^n(u) \vec{1}$$

where $\mathbf{b}^t(u)$ is an initial vector and $\vec{1} = (1, \dots, 1)$.

7. Spectral Decomposition

Let $\lambda(u)$ be the largest eigenvalue of $\mathbf{T}(u)$. Then

$$O_n(u) = c(u) \lambda^n(u) (1 + O(A^n))$$

for some $A < 1$. This proves that the generating function $O_n(u)$ satisfies the so called quasi-power law.

8. The above formula suggests that we deal with weakly dependent random variables described by the generating function $\lambda(u)$. Hence, for example

$$\mathbf{E}[O_n] = n\lambda'(1) + O(1).$$

Final Results

Mean and Variance

$$\begin{aligned}\mathbf{E}[O_n(\mathcal{W})] &= n\Lambda'(0) + O(1) = nP(\mathcal{W}) + O(1), \\ \mathbf{Var}[O_n(\mathcal{W})] &= n\Lambda''(0) + O(1) = n\sigma^2(\mathcal{W}) + O(1)\end{aligned}$$

where $\Lambda(s) = \log \lambda(e^s)$

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Central Limit Theorem

$$\Pr \left\{ \frac{O_n - nP(\mathcal{W})}{\sigma(\mathcal{W})\sqrt{n}} \leq x \right\} \sim \frac{1}{\sqrt{2\pi}} \int_{-\infty}^x e^{-t^2/2}$$

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Large Deviations

If $\mathbf{T}(u)$ is **primitive**, then

$$\Pr\{O_n(\mathcal{W}) = a\mathbf{E}[O_n]\} \sim \frac{1}{\sigma_a\sqrt{2\pi n}} e^{-nI(a)+\theta_a}$$

where $I(a)$ can be explicitly computed, and θ_a is a known constant.

Based on: [Flajolet, W.S.](#) and [Vallee](#), JACM, 2006.

Some Experiments

The complete works of Shakespeare are found under

<http://the-tech.mit.edu/Shakespeare/>.

We first extracted the full text of Hamlet stripped of all the comments and searched for

$\mathcal{W} = \text{thelawisgaussian}$

		$w = \text{thelawisgaussian}$		$\tilde{w} = \text{naissuagsiwaleht}$	
d	Expected (E)	Occurred (O_n)	O_n/E	Occurred (O_n)	O_n/E
13	9.195E+01	0	0.00	18	0.19
14	2.794E+02	693	2.47	371	1.32
15	7.866E+02	1,526	5.46	2,379	3.02
18	1.211E+04	31,385	2.58	14,123	1.16
20	5.886E+04	124,499	2.11	41,066	0.69
25	1.673E+06	2,527,148	1.51	1,277,584	0.76
30	2.577E+07	40,001,940	1.55	25,631,589	0.99
40	1.928E+09	2,757,171,648	1.42	2,144,491,367	1.11
50	5.482E+10	76,146,232,395	1.38	48,386,404,680	0.88
∞	1.330E+48	1.36554E+48	1.03	1.38807E+48	1.04

Figure 3: Observed occurrences (O_n) versus predicted values (expectations, E) in the alphabetical characters of Hamlet.

Reliable Threshold for Intrusion Detection

We argued that one needs a reliable threshold for intrusion detection. If false alarms are to be avoided, the problem is of finding a threshold $\alpha_0 = \alpha_0(\mathcal{W}; n, \beta)$ such that

$$P(O_n(\mathcal{W}) > \alpha_{th}) \leq \beta (= 10^{-5}).$$

Our results shows that

$$\alpha_{th} = nP(\mathcal{W}) + x_0\sigma(\mathcal{W})\sqrt{n}, \quad \beta = \frac{1}{\sqrt{2\pi}} \int_{x_0}^{\infty} e^{-t^2/2} dt \sim \frac{1}{x_0} e^{-x_0^2/2}.$$

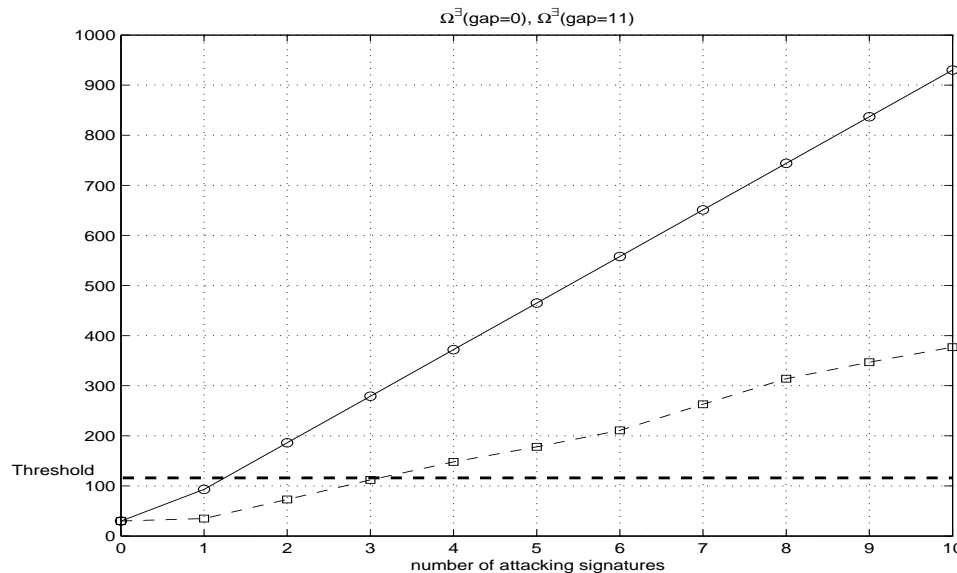


Figure 4: Pattern=wojciech, window=100 (cf. Gwadrea et al. (2004)).

Outline Update

1. Motivations
2. Pattern Matching Problems
3. Analysis & Applications
 - Exact String Matching & Finding Biological Motifs
 - Hidden Patterns & Intrusion Detection
 - Joint String Complexity & Classification of Twitter

Some Definitions

String Complexity of a single sequence is the number of **distinct** substrings.

Throughout, we write X for the string and denote by $I(X)$ the set of *distinct substrings* of X over alphabet \mathcal{A} .

Example. If $X = abaa$, then

$$I(X) = \{\epsilon, a, b, aa, ab, ba, aab, aba, baa, aaba, abaa, aabaa\},$$

so $|I(X)| = 12$. But if $X = aaaaa$, then

$$I(X) = \{\epsilon, a, aa, aaa, aaaa, aaaaa\},$$

so $|I(X)| = 6$.

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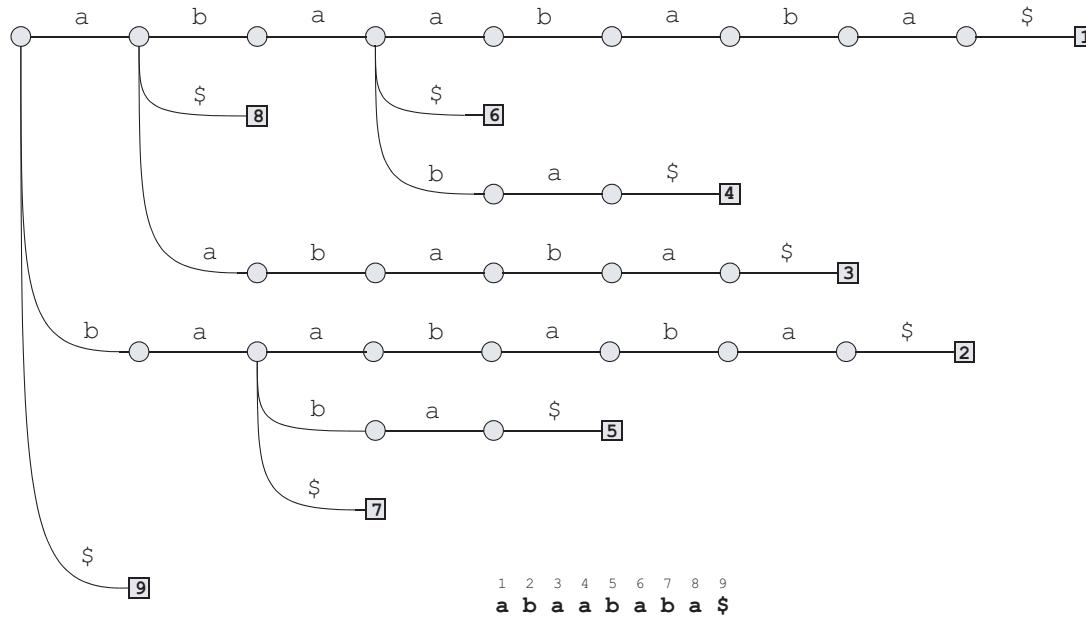
so $|I(X)| = 6$.

The **string complexity** is the **cardinality** of $I(X)$ and we study here the *average* string complexity

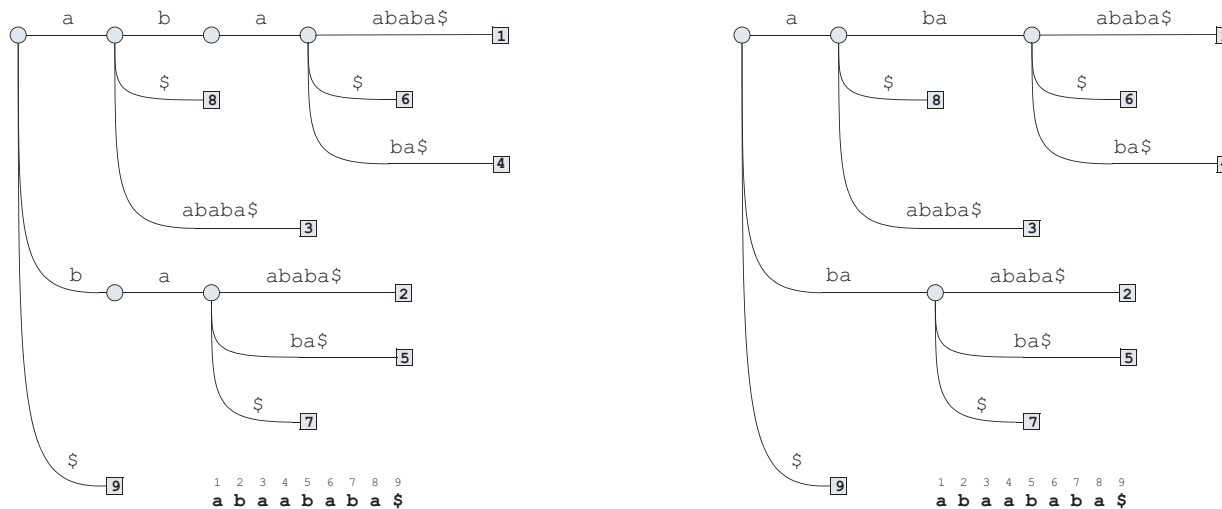
$$\mathbf{E}[|I(X)|] = \sum_{X \in \mathcal{A}^n} P(X) |I(X)|.$$

where X is generated by a **memoryless/Markov** source.

Suffix Trees and String Complexity



Non-compact suffix trie for $X = \text{abaababa}$ and string complexity $I(X) = 24$.



String Complexity = # internal nodes in a non-compact suffix tree.

Some Simple Facts

Let $O(w)$ denote the number of times that the word w occurs in X . Then

$$|I(X)| = \sum_{w \in \mathcal{A}^*} \min\{1, O(w)\}.$$

Since between any two positions in X there is one and only one substring:

$$\sum_{w \in \mathcal{A}^*} O(w) = \frac{(|X| + 1)|X|}{2}.$$

Hence

$$|I(X)| = \frac{(|X| + 1)|X|}{2} - \sum_{w \in \mathcal{A}^*} \max\{0, O(w) - 1\}.$$

Define: $C_n := \mathbf{E}[|I(X)| \mid |X| = n]$. Then

$$C_n = \frac{(n + 1)n}{2} - \sum_{w \in \mathcal{A}^*} \sum_{k \geq 2} (k - 1)P(O_n(w) = k).$$

We need to study probabilistically $O_n(w)$: that is:

number of w occurrences in a text X generated a probabilistic source.

Methodology and Some Results

Last expression allows us to write

$$C_n = \frac{(n+1)n}{2} + \mathbf{E}[S_n] - \mathbf{E}[L_n]$$

where $\mathbf{E}[S_n]$ and $\mathbf{E}[L_n]$ are, respectively, the average size and path length in the associated (compact) suffix trees.

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We know that (Jacquet & Regnier, 1989; W.S., 2001)

$$\mathbf{E}[S_n] = \frac{1}{h}(n + \Psi(\log n)) + o(n), \quad \mathbf{E}[L_n] = \frac{n \log n}{h} + n\Psi_2(\log n) + o(n),$$

where $\Psi(\log n)$ and $\Psi_2(\log n)$ are periodic functions. Therefore,

$$C_n = \frac{(n+1)n}{2} - \frac{n}{h}(\log n - 1 + Q_0(\log n) + o(1))$$

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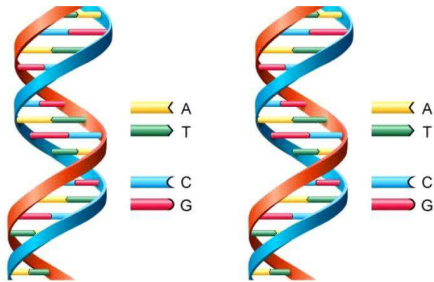
Theorem 2 (Janson, Lonardi, W.S., 2004). For unbiased memoryless source:

$$C_n = \binom{n+1}{2} - n \log_{|\mathcal{A}|} n + \left(\frac{1}{2} + \frac{1-\gamma}{\ln |\mathcal{A}|} + Q_1(\log_{|\mathcal{A}|} n) \right) n + O(\sqrt{n \log n})$$

where $\gamma \approx 0.577$ is Euler's constant and Q_1 is a periodic function.

Joint String Complexity

For X and Y , let $J(X, Y)$ be the set of **common words** between X and Y .



The joint string complexity is

$$|J(X, Y)| = |I(X) \cap I(Y)|$$

Example. If $X = aabaa$ and $Y = abbba$, then $J(X, Y) = \{\varepsilon, a, b, ab, ba\}$.

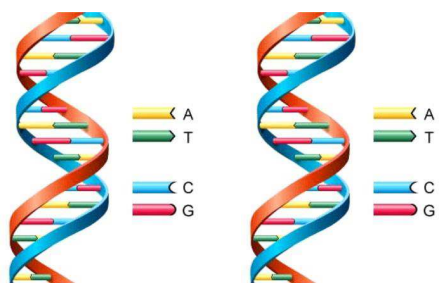
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when $|X| = n$ and $|Y| = m$.

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Some Observations. For any word $w \in \mathcal{A}^*$

$$|J(X, Y)| = \sum_{w \in \mathcal{A}^*} \min\{1, O_X(w)\} \cdot \min\{1, O_Y(w)\}.$$

When $|X| = n$ and $|Y| = m$, we have

$$J_{n,m} = \mathbf{E}[|J(X, Y)|] - 1 = \sum_{w \in \mathcal{A}^* - \{\varepsilon\}} P(O_n^1(w) \geq 1) P(O_m^2(w) \geq 1)$$

where $O_n^i(w)$ is the number of w -occurrences in a string of generated by source $i = 1, 2$ (i.e., X and Y) which we assume to be **memoryless sources**.

Independent Joint String Complexity

Consider two sets of n independently generated (memoryless) strings.

Let $\Omega_n^i(w)$ be the number of strings for which w is a prefix when the n strings are generated by a source $i = 1, 2$ define

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for large n .

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Recurrence for $C_{n,m}$

$$C_{n,m} = 1 + \sum_{a \in \mathcal{A}} \sum_{k, \ell \geq 0} \binom{n}{k} P_1(a)^k (1 - P_1(a))^{n-k} \binom{m}{\ell} P_2(a)^\ell (1 - P_2(a))^{m-\ell} C_{k,\ell}$$

with $C_{0,m} = C_{n,0} = 0$.

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Main Results

Assume that $\forall a \in \mathcal{A}$ we have $P_1(a) = P_2(a) = p_a$.

Theorem 4. For a *biased memoryless source*, the *joint complexity* is asymptotically

$$C_{n,n} = n \frac{2 \log 2}{h} + Q(\log n)n + o(n),$$

where $Q(x)$ is a small *periodic function* (with amplitude smaller than 10^{-6}) which is *nonzero* only when the $\log p_a$, $a \in \mathcal{A}$, are *rationally related*, that is, $\log p_a / \log p_b \in \mathbb{Q}$.

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Assume that $P_1(a) \neq P_2(a)$.

Theorem 5. Define $\kappa = \min_{(s_1, s_2) \in \mathcal{K} \cap \mathbb{R}^2} \{(-s_1 - s_2)\} < 1$, where s_1 and s_2 are *roots* of

$$H(s_1, s_2) = 1 - \sum_{a \in \mathcal{A}} (P_1(a))^{-s_1} (P_2(a))^{-s_2} = 0.$$

Then

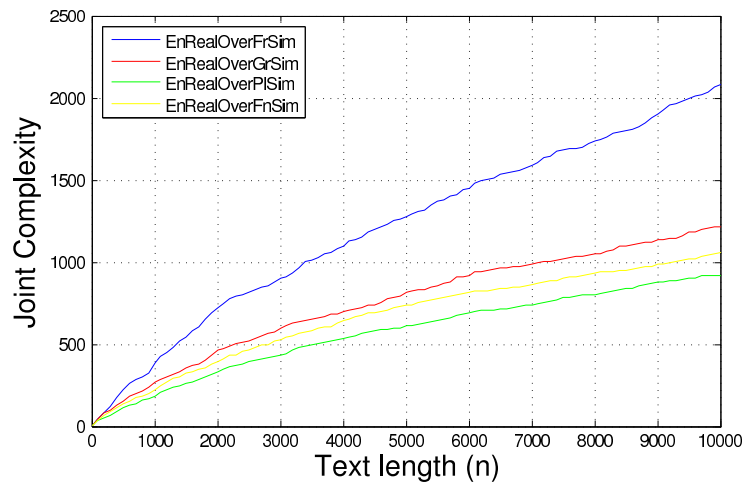
$$C_{n,n} = \frac{n^\kappa}{\sqrt{\log n}} \left(\frac{\Gamma(c_1)\Gamma(c_2)}{\sqrt{\pi \Delta H(c_1, c_2) \nabla H(c_1, c_2)}} + Q(\log n) + O(1/\log n) \right),$$

where Q is a double periodic function.

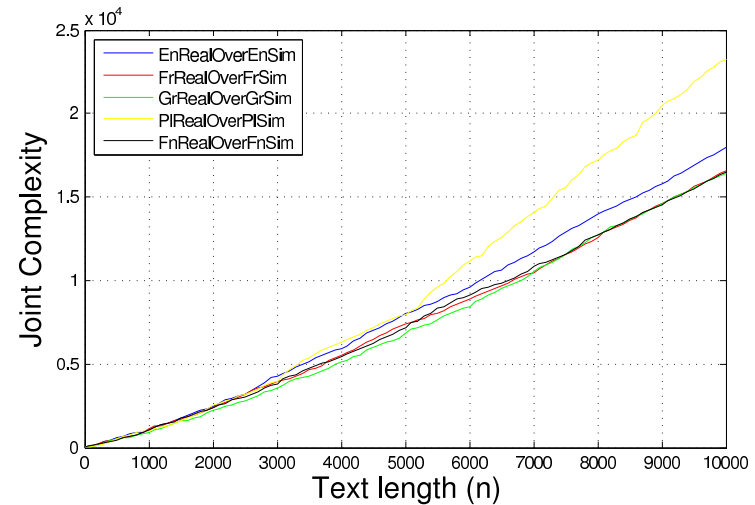
Classification of Sources

The growth of $C_{n,n}$ is:

- $\Theta(n)$ for **identical sources**;
- $\Theta(n^\kappa / \sqrt{\log n})$ for **non identical sources** with $\kappa < 1$.



(a)



(b)

Figure 5: Joint complexity: (a) English text vs French, Greek, Polish, and Finnish texts; (b) real and simulated texts (3rd Markov order) of English, French, Greek, Polish and Finnish language.

Acknowledgments

My French Connection:



Philippe Flajolet (1948-2011)

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. . . and **ALL** my co-authors.

That's It



THANK YOU