

Pattern matching

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Pattern matching

- Important and classical question in computer science
- In practice it arises in many application contexts bioinformatics, text processing, databases, etc.
- Different formulations depending on whether one searches for
 - ① one or several words
 - ② exactly or approximately
 - ③ regular expression
 - ④ set of similar words

Different algorithmic approaches to exact pattern matching

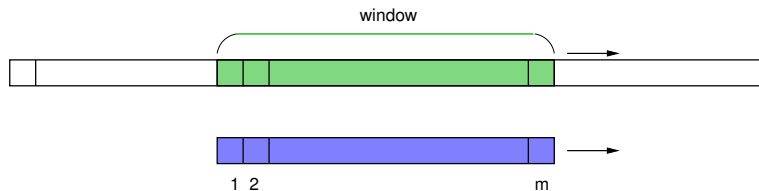
- Automaton based
e.g., [Aho Corasick 75]
- Window scan algorithms
e.g., [Horspool 80]
- Algorithms using bit parallelism
e.g., [Baeza-Yates Perleberg 86]
- Fingerprints
e.g., [Karp Rabin 87]

From simple to complex: Different types of patterns

Exact Pattern Matching

- 1 a text T of length n
- 2 a pattern M of length m , and generally $m \ll n$.

For single word: [window scan algorithm](#)



Exact Pattern Matching

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Example: $M := tgtg$

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
T:	c	t	g	t	g	t	g	t	a	c	a	t	g	t	g

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Solution: $\{2, 4, 12\}$

Exact Set Pattern Matching

Search simultaneously for occurrences of a **set of words** in a text.

Input: a set $\mathcal{M} := \{tgtg, atg, cat\}$ of words and a text $T := ctgtgtgtacatgtg$.

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M_1 at positions $\{2, 4, 12\}$

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Solutions:

M_1 at positions $\{2, 4, 12\}$

M_2 at position $\{11\}$

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Solutions:

M_1 at positions $\{2, 4, 12\}$

M_2 at position $\{11\}$

M_3 at position $\{10\}$

Approximate Pattern Matching

- 1 Idem: a *text* T of length n , a *pattern* M of length m
- 2 a maximum number of allowed differences.

Example: $M := tgtg$

One mismatch allowed (at most)

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						t	g	t	g						

Additional occurrence with one mismatch at pos. 6 in T
mismatch at pos. 4 in M

Solution: $\{2, 4, 6, 12\}$

Probabilistic motif search: Position Weight Matrix (PWM)

Definition: Position Weight Matrix (PWM)

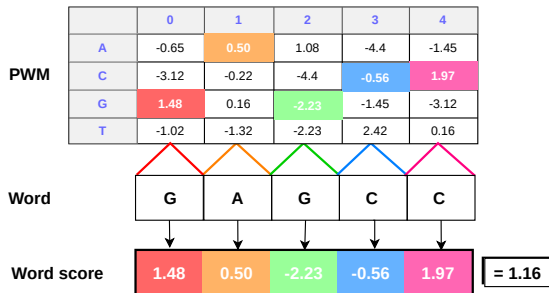
For DNA, a PWM M is $4 \times m$ matrix:

- for $\alpha \in \Sigma$ and position i , entry $M[\alpha, i] :=$ the score of nuc. α at position i
- The score of a word is the sum of scores at all positions.

Problem:

Given a text T , a PWM M , a score threshold s , find all substrings of T whose score is $> s$.

Scoring of a
word
(or a substring
of T)



Searching for probabilistic motifs

PWMs are the simplest model

More complex motifs exist

Searching for probabilistic motifs

PWMs are the simplest model

dinucleotidic PWMs
HOCOMOCO database
for Transcription Factors

More complex motifs exist

di-PWM

	0	1	2	3	4
AA	0.77	-0.14	-0.14	-1.97	-1.77
AC	-0.78	-1.97	-2.58	-2.58	-2.23
AG	-0.65	0.50	1.08	-4.4	-1.45
AT	-1.77	-1.2	-4.4	1.27	-4.4
CA	0.52	0.26	-1.11	-3.18	0.64
CC	-1.77	-0.43	-3.12	-0.56	-1.6
CG	-1.32	0.94	0.31	-4.4	-0.14
CT	-3.12	-0.22	-4.4	-1.77	1.97
GA	1.48	0.16	0.82	-1.45	-3.12
GC	0.33	-0.43	-2.23	-1.97	-4.4
GG	1.28	0.85	1.83	-1.97	-1.97
GT	-1.02	-1.32	-2.23	2.42	0.16
TA	-1.21	-0.59	-1.11	-4.4	0.61
TC	-2.23	-1.11	-4.4	-4.4	2.14
TG	-1.45	0.78	0.19	-4.4	-0.54
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di-PWM

dinucleotidic PWMs

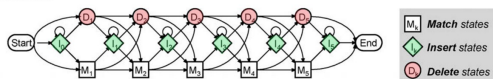
HOCOMOCO database
for Transcription Factors

profile Hidden

Markov Models (HMMs)

PFAM database

Profile-HMM



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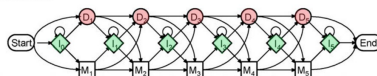
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Other motif representations: gapped motifs, Covariance Matrix [Durbin et al. 98].

Exact pattern matching: a primer

- 1 The problem
- 2 Naive algorithm
- 3 Linear time algorithms
- 4 Text indexing approach
- 5 Filtration approach

Exact Pattern Matching

b b a b a c a c a a c a b a b a a b b a b

Text T of length t

Exact Pattern Matching

b b a b a c a c a a c a b a b a a b b a b

Text T of length t

b a b a

Word M of length m

Exact Pattern Matching

Position of
occurrences?

b b a b a c a c a a c a b a b a a b b a b

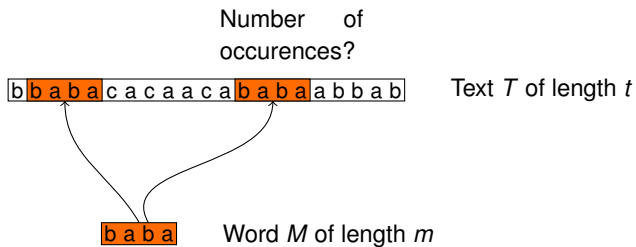
Text T of length t

b a b a

Word M of length m



Exact Pattern Matching



Naive and involved algorithms

- **Naive** algorithm:
for each window m pairwise symbol comparisons
about n windows
Total **time** proportional to $n * m$ (complexity)
- **Linear time** solutions:
Idea: exploit results on a window to ease that of overlapping windows
Boyer-Moore or Knuth Morris Pratt algorithms in the 70's
Total time proportional to $n + m$

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Limitations: single query and exact match
Answers: indexing text and filtration approaches

Naive algorithm : scan and shift

- basic operation: char. equality test $O(1)$
- compare M to a window of size m testing for char equality
- repeat for all $(n - m)$ possible windows
- time complexity $O((n - m) \times m)$
- *scanning* direction in T : usually left to right
- *verification* of current window: left to right
- *shift*: distance between two windows considered successively
- naive algorithm: shifts equal one, which is minimum

Forward search: Morris Pratt

- [Morris, Pratt, 76]
- both scanning and verification from left to right
- *safe shift*: a shift that do not skip over potentially valid windows
- verification: prefix wise, from left to right
- scanning left to right
- shift use borders of prefixes of M
- Later improved by Knuth into Knuth-Morris-Pratt algorithm [Knuth, Morris, Pratt, 77].

Algorithme 1 : Morris-Pratt

Input : Text T of length n and pattern M of length m

Precompute the Shift table of M ;

$i := 1; j := 1;$

while ($i < n - m$) **do**

$j := 0;$

while ($j < m$) *et* ($M[j] = T[i + j]$) **do**

$j := j + 1;$

if ($j = m$) **then**

 print (occurrence of M at position i);

$i := i + \text{Shift}[j];$

$j := \max(0, j - \text{Shift}[j]);$

Backward search: Horspool

- Simplification of Boyer Moore algorithm
Verification from right to left – [Horspool, 80]
- uses a single rule for shifts: improved *Bad Character rule*
considers only the last symbol in the current window

Complexity - efficiency

- Boyer Moore worst case time complexity is $O(n + m)$
- Horspool: worst case $O(m \times n)$!
- However efficient in practice: sublinear expected running time

Horspool Algorithm

i : offset for index in pattern and in text, j : index text current window

Algorithm 2 : Horspool

Input : Text T of length n and pattern M of length m

Precompute L the shift table of M ;

$j := 0$;

while ($j < n - m$) **do**

$i := m - 1$;

while ($i > 0$) *et* ($M[i + 1] = T[j + i + 1]$) **do**

$i := i - 1$;

if ($i \leq 0$) **then**

 print (occurrence of M at position $j + 1$);

$j := j + m - L[T[j + m]]$;

Shift table L for pattern M

for a symbol c : gives the position of the rightmost occurrence of c in $M[1; m-1]$

Algorithm 3 : Horspool preprocessing

Input : Pattern M of length m

Output: returns table L of length σ

Precompute L the shift table of M ;

for $c \in \Sigma$ **do**

$L[c] := 0$;

for $i := 1..m-1$ **do**

$L[M[i]] := i$;

Set Pattern Matching: searching for several motifs

Multiple motifs search

Given

- $\mathcal{M} = \{M_1, \dots, M_z\}$ a set of z words of length m_1, \dots, m_z ,
- T a text of length n .

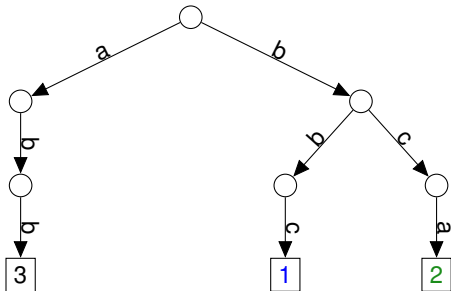
Question: find all occurrences of each motif in T .

- **Aho-Corasick** (AC) algorithm: uses a tree to represent the motifs
- **Motifs trie:** each branch of the tree spells out one motif
Failure links: links a suffix of a window to the largest prefix of some M_i
preprocessing is linear in the sum of the motif lengths.
- Naive Algorithm: scan the text T using the tree without failure links
takes quadratic time.
- AC algorithm scans the text T only once, takes linear time $O(n)$. [Aho Corasick 75]

Aho Corasick automaton: example 1

Let $\mathcal{M} = \{abb, bbc, bca\}$

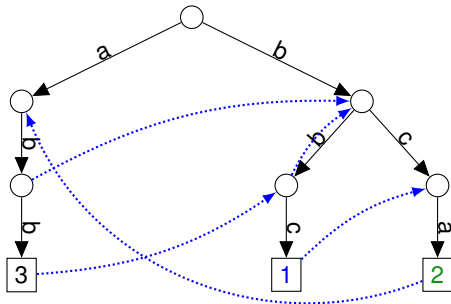
Trie of words in \mathcal{M}
with failure links as dotted blue lines.



Aho Corasick automaton: example 1

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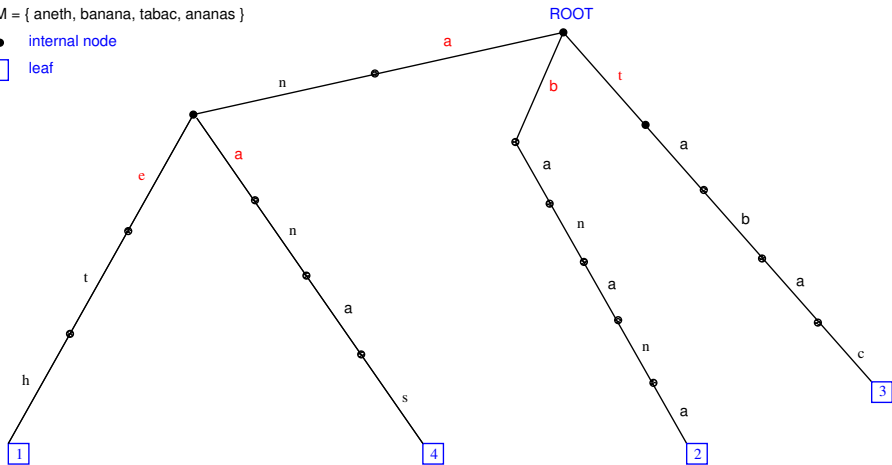


Example 2

$M = \{ \text{aneth, banana, tabac, ananas} \}$

● internal node

□ leaf



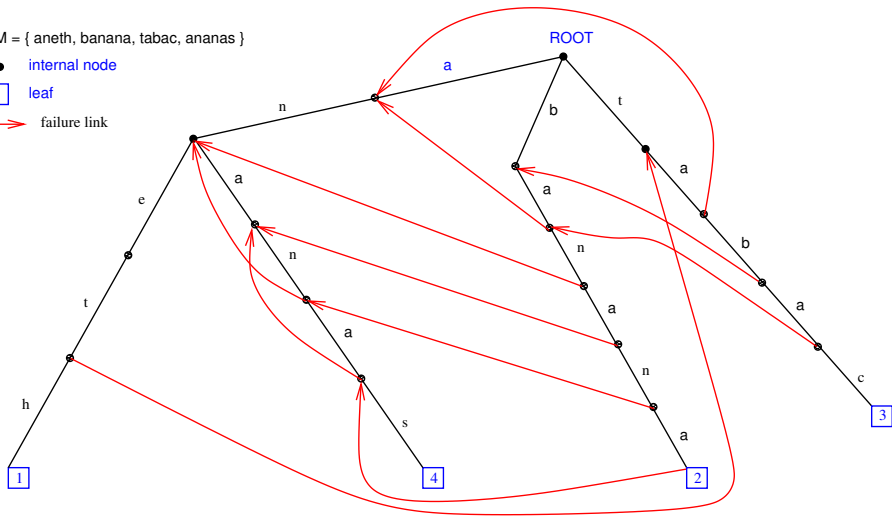
Example 2

$M = \{ \text{aneth, banana, tabac, ananas} \}$

● internal node

□ leaf

→ failure link



Off-line search: indexing the text for optimal search time

Matching in two steps:

- 1 **preprocessing** the text T in time $O(n)$
build and store a data structure: an **index**
enables exact search query
- 2 **search** for each pattern in the index in $O(m)$ time (optimal)

Text indexing data structures

For a text of length n , a **good** index:

- 1 occupancy memory in $O(n)$
- 2 construction time in $O(n)$ units
- 3 enables exact motif search in $O(m)$ time for a motif of length m

Three historical structures:

- 1 **compact suffix tree** [Wiener 73, McCreight 76, Ukkonen 92]
- 2 **suffix array**: construction in $O(n)$ [Manber & Myers 90, Kärkkäinen & Sanders 03]
- 3 **DAWG** (Directed Acyclic Word Graph) [Blumer et al. 85]

Breakthrough in text indexing

With historical index structures,

- 1 you need the text and the index
- 2 both in main memory to keep it fast

Around 2000, the advent of compressible “self indexing structures”:

- 1 a self-index replaces the text and the classical index
- 2 its size can be modulated in function of available memory.

Example

- 1 Burrows-Wheeler Transform or FM-index [Ferragina Manzini 00]
- 2 Enhanced Suffix Arrays [Ohlebusch, 13]
- 3 Various compressed k -mer indexes

Used in other contexts: overlap graphs, de Bruijn Graphs construction.

PM using bit parallelism

PM using state array and binary operations

- Often $|M| \ll |T|$ and M fits into a machine word
- safely shifting the window requires knowing which prefixes of M match the current window
- store this information in a state array
- shift is performed using binary operation

Avantage: fast

binary operations \Rightarrow time efficiency

Example

Shift-OR [Baeza-Yates & Perleberg, 96]

Variation of Shift-And algorithm [Baeza-Yates & Gonnet, 92]

Binary operations

- unsigned integers (int): as binary vectors with 16 or 32 bits
- unsigned short int: 16 bits
- unsigned long int: 32 bits
- binary encoding from right to left

Examples of binary encodings

value	encoding on 16 bits
0	0000000000000000
1	0000000000000001
2	0000000000000010
3	0000000000000011
4	0000000000000100
2^{i-1}	1 at i^{th} position from right

Binary operations

- \ll : shift to the left
- \gg : shift to the right
- $\&$: binary AND
- $|$: binary OR
- \wedge : binary XOR
- \sim : binary complement (negation)

Example of operations

value	encoding
~ 0	1111111111111111
~ 3	1111111111111100
$1 \ll 1$	0000000000000010
$5 \gg 2$	0000000000000001

Shift-OR: principle

- a pattern M and a text T
- current window ends at position p in T
- State array E : a binary array of length m , as many as prefixes of M
- E_p encodes which prefixes of M are also suffixes of current window ending at pos. p
- if the bit corresponding to entire M equals zero \Rightarrow occurrence of M
- using $T[p+1]$ and E_p , one easily computes E_{p+1}

Illustration of state array (1)

Consider

- a word $M = \text{acat}$ of length 4

- a text $T = \text{gacat}$ of length 5

T contains 2 windows of length 4: *gaca* puis *acat*

Illustration of state array (2)

The state array E

at position 4

Illustration of state array (2)

The state array E

1	2	3	4	E_4
g	a	c	a	

at position 4

Illustration of state array (2)

The state array E

1	2	3	4	E_4
g	a	c	a	
			a	0

at position 4

Illustration of state array (2)

The state array E

1	2	3	4	E_4
g	a	c	a	
			a	0
		\neq		1

at position 4

Illustration of state array (2)

The state array E

1	2	3	4	E_4
g	a	c	a	
			a	0
		\neq		1
	a	c	a	0

at position 4

Illustration of state array (2)

The state array E

1	2	3	4	E_4
g	a	c	a	
			a	0
		\neq		1
	a	c	a	0
\neq				1

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2	3	4	5	E_5
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≠				1

then at position 5

2	3	4	5	E_5
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		a	≠	1
	≠			1
a	c	a	t	0

Question: how to compute E_5 from E_4 and $T[p+1]$?

Decomposing equality of prefix of M and a suffix of current window in T

Let c and p be two integers such that

- $1 < c \leq m$ and
- $0 < p \leq n$.

We get $M[1, c] = T[p - c + 1, p]$

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$$M[1, c - 1] = T[p - c + 1, p - 1] \text{ and } T[p] \text{ matches the } c^{\text{th}} \text{ position of } M$$

Shift-OR: bit masks

Idea: for each symbol α , a bit mask indicates the positions of α in M .

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a	$L_a = 1010$

$M = acat$

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Example

	letter	bit mask (lightest bit on right)
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	c	$L_c = 1101$
	g	$L_g = 1111$
	t	$L_t = 0111$

Shift-OR principe: final

Let E_p be the state array of m bits such that for all $j \in [1, m]$ one has:

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Thanks to the decomposition, one gets:

$$\begin{aligned} E_p[c] &= E_{p-1}[c-1] \text{ OR } L_{T[p]}[c] \quad \text{if } c > 1 \text{ AND} \\ E_p[1] &= L_{T[p]}[1] \end{aligned}$$

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Idea: thanks to bit arrays, one compute all bits in parallel (with binary operations)

$$E_p := (E_{p-1} \ll 1) \text{ OR } L_{T[p]}.$$

Algorithme 4 : Shift-OR

Input : Text T et pattern M resp. of lengths n and m

Preprocessing: compute bit masks in L ;

$E := \underbrace{1\dots 1}_{m \text{ times}};$

for p from 1 to n **do**

$E := (E \ll 1) \text{ or } L_{T[p]};$

si $(E < 2^{m-1})$ **alors**

 report an occurrence of M ending at position p in T ;

Probability of occurrence

Consider all words of length q : the set $\Sigma^q := \{Q_i : 0 < i \leq \sigma^q\}$.

Denote by $\Pr(Q_i \notin T)$: the **absence probability** of a word in a random text T .
i.e., the fact that Q_i does not occur in T .

- The probability $\Pr(Q_i \notin T)$ is not the same for all Q_i .
It depends on the form of Q_i .
More precisely: On the *autocorrelation* of Q_i .
- **Autocorrelation**: binary vector storing the “Periodicity” of Q_i i.e., the set of periods.

Definition (Period)

Let $Q \in \Sigma^q$ and let p be a non-negative integer with $p < q$.
Then p is a **period** of Q iff:

$$\forall 0 \leq i < n - q : Q[i] = Q[i + p].$$

Example

Shift:	0	1	2	3	4	5	6	7	8	9	0		
Word:	A	B	R	A	C	A	D	A	B	R	A	:	:

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Word:	A	B	R	A	C	A	D	A	B	R	A	:	:						
							A	B	R	A	C	A	D	A	B	R	A	:	:

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						A	B	R	A	C	A	D	A	B	R	A	:	:
							A	B	R	A	C	A	D	A	B	R	A	

Period set of *ABRACADABRA* = $\{0, 7, 10\}$.

Autocorrelation : example

- $Q=ABRACADABRA$. **M0 model** for random texts.

$$P(A) = 0.4, P(B) = 0.2, P(C) = 0.1, P(D) = 0.1, P(R) = 0.2.$$

- How does Q overlap with itself?

Shift:	0	1	2	3	4	5	6	7	8	9	0	$P(\text{Tail})$
ABRACADABRA												$P(\epsilon) = 1$
												$P(\text{CADABRA}) = \frac{256}{10^7}$
												$P(\text{BRACADABRA}) = \frac{4096}{10^{10}}$

- **Autocorrelation vector:**

$$A_Q = (1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1)$$

- **Autocorrelation polynomial:**

$$C_Q(z) := \sum_{j=0}^{q-1} A_Q(j) z^j = 1 + \frac{256}{10^7} z^7 + \frac{4096}{10^{10}} z^{10}$$

P(word Q does not occur in text T)

- Word Q with autocorrelation polynomial $C(z)$
Text $T^{(n)}$ of length n
- **Theorem:** Generating function in M0-model:

$$P(z) = \frac{C(z)}{\Pr(Q)z^q + (1-z) \cdot C(z)} = \sum_{j \geq 0} p_j z^j$$

[Guibas & Odlyzko 81a,81b, Chrysaphinou & Papastavridis 91]

- When $P(z)$ is rational, in certain conditions, one can compute p_n , i.e., the coefficient of the $(n+1)^{th}$ term of $P(z)$, that is the probability that Q does not occur in $T^{(n)}$.

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Extensions to estimate **how many q-grams**

- 1 are **missing** from a random text
- 2 are **common** to two random texts.

[Rahmann Rivals 00 & 03]

Approximate Pattern Matching

Approximate Pattern Matching

Given:

- a pattern M of m characters
- a text T of n characters
- k an integer such that $k \leq m$
- a cost function $e(U, V)$ between two strings U, V the unit cost edit distance.

Definition : find the starting positions i in T of any substring P such that $e(M, P) \leq k$.

Complexity : worst case $O(kn)$ with $O(m^2)$ preprocessing of P .

Others :
algorithm with sublinear expected time [Chang & Lawler, 1990]
filtration algorithms are efficient in practice.

Two phases algorithm: **filtration** and **checking**
based on a necessary condition (NC) for a match

Filtration : find all substrings P' of T verifying the NC
 P' is called a *potential match*

Checking : check if a potential match is a match
this for all potential matches, use dynamic programming in $O(nm)$ time

Advantage

if the NC is easy to compute and potential matches are seldom,
only few substrings are checked using dynamic programming algorithm
⇒ Gain of execution time

- Reduction to exact partitionning [Baeza-Yates, Perleberg, 92]
- Maximal matches distance [Chang, Lawler, 90] [Ukkonen, 92]
- q -gram distance [Owolabi, Mc Gregor, 88], [Jokinen, Ukkonen, 91]
- Double filtration with gapped tuple (Pevzner-Waterman 94)

Heuristics: BLAST, FASTA, d2 [Torney et al, 90]

Idea

- 1 cut the pattern in $k + 1$ adjacent substrings of length $\lfloor \frac{m}{k+1} \rfloor$
- 2 search for all pieces
- 3 if at most k errors are allowed, at least one piece matches exactly

Generalisations : a) cut in $k + s$ pieces and search for s distinct pieces conserving order in the pattern,
b) cut in j pieces and search each piece with $\lfloor \frac{k}{j} \rfloor$ errors

With index : [Baeza-Yates Navarro 96] use a table of the q -grams occurrences and reduce pieces to $q - gram$

Definition: q -gram or q -mer

A q -gram is a string of length q over an alphabet Σ .

Idea

Let $q \leq \lfloor \frac{m}{k+1} \rfloor$.

- count the number of matching q -grams between P and P'
- when $e(P, P') \leq k$, each error kills at most q q -grams
- thus at least $m - (k+1)q + 1$ q -grams match between P and P' .

Definition : a q -gram is a string of length q over an alphabet Σ

Idea : Let $q \leq \lfloor \frac{m}{k+1} \rfloor$, count the nb of q -grams equal between M & M' si $e(M, M') \leq k$, each difference affect at most q q -grams.

Worst case : $m - (k + 1)q + 1$ q -grams match between M & M' .

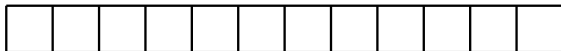
q -gram filter

length of M : 12; $q := 4$

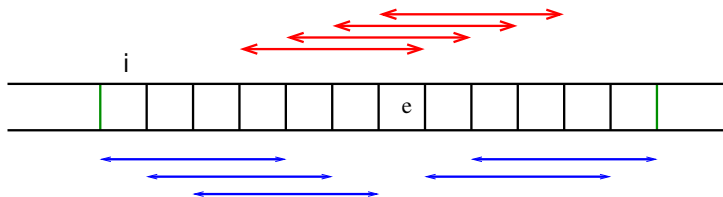
↔: equal q -grams

↔: different q -grams

M



T



[Owolabi, McGregor, 88]



Thanks for your attention

Questions?



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