# Fast nGram-Based String Search Over Data Encoded Using Algebraic Signatures 

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## Plan

- Problem Statement
- Our Proposal
- Key Idea
$\square$ Algebraic Signatures
$\square$ Record Encoding
$\square$ Pattern Preprocessing
- Search Example
- Performance Study
- Conclusion


## Problem

- String Search (Pattern Matching) in A Database or File
$\square$ Find every record matching pattern = "Dauphine"
$\square$ What about record "Universite de Technologie Paris Dauphine" ?
- Records are searched often, and updated rarely
■ We especially target large Scalable and Distributed DBs and Files
- on Grids and P2P networks



## Our Proposal

- Fast String Search Method
$\square$ Several Times Faster than Boyer-Moore
- In our experiments:
$\square$ Up to eleven times for ASCII
$\square$ Up to six times for XML
$\square$ Up to seventy times for DNA


## Key Idea : Pre-processing

■ We aggregate (encode) all $n$-symbol long substrings (ngrams) in visited strings (records) and in the searched pattern into single-symbol algebraic signatures
$\square$ Records are encoded while coming for storage
$\square$ Pattern is encoded during search preprocessing


## Key Idea : Search

- We compare signatures for attempted matches and shifts like Boyer-Moore (BM) does
- "Bad character" shift
- However, matching ngram signatures $\Leftrightarrow$ matching $n$ symbols at the time


## Key Benefit

- Matching attempts usually more discriminative than matching a single (original) symbol at the time.
$\square$ The latter is the current approach
- BM and all other major pattern matching algorithms we are aware of
$\square$ KMP, Quick Search, KR...


## Key Benefit

- Longer shifts
- Fewer comparisons
- Faster search
- Local search over encoded data only
- No local user can claim unintentional disclosure of stored data
- Important for P2P
- Thought determined fraud is not that difficult
- Idem for the data transfer to the client


## Algebraic Signature ICDE 2004

- Condenses information in a string into a single character
- Defined over Galois Fields (GF) of size $2^{f}$
$\square$ Elements are bit strings of length $f$
$\square$ In our case, typically $f=8$
$\square$ Hence our symbols are bytes
$\square$ We realize GF addition $\oplus$ as XOR
$\square$ We realize GF multiplication through log/antilog tables


## Algebraic Signature

$$
A S\left(r_{1} \ldots r_{k}\right)=r_{1} \alpha \oplus r_{2} \alpha^{2} \oplus \cdots \oplus r_{k} \alpha^{k}
$$

$\Rightarrow \alpha$ is a primitive element, e.g., $\alpha=2$
$\Rightarrow$ if $A S\left(R_{1}\right) \neq A S\left(R_{2}\right)$ then $R_{1} \neq R_{2}$ for sure
$\Rightarrow$ if $A S\left(R_{1}\right)=A S\left(R_{2}\right)$ then for sure or very likely $R_{1}=R_{2}$
$\square$ The latter case is a collision

## Record Encoding

- We encode every stored record : $r_{1} \ldots r_{K}$
$\square$ Either into full Cumulative Algebraic Signature

$$
r_{k}^{\prime}=r_{1} \alpha \oplus r_{2} \alpha^{2} \oplus \cdots \oplus r_{k} \alpha^{k}
$$

$\square$ Or into partial (moving) CAS of ngrams
$r_{k}^{\prime}=r_{k-n+1} \alpha \oplus \cdots \oplus r_{k} \alpha^{n}$

## Full CAS




## Partial CAS for $n=2$



- Partial CAS can be stored or dynamically calculated from full CAS
- See the paper


## Pattern Preprocessing

- We aggregate ngram signatures in the pattern in a BM-like shift table T
- Conceptual result for "Dauphine"
- Actually:
$\square$ shift table size is $f$ and entry is by AS value
$\square$ Rightmost ngram value is in variable $V$

| 2-gram | Shift |
| ---: | ---: |
| $33=$ AS (da) | 6 |
| $23=$ AS (au) | 5 |
| $133=$ AS (up) | 4 |
| $24=$ AS (ph) | 3 |
| $07=$ AS (hi) | 2 |
| $62=$ AS (in) | 1 |
| $67=$ AS (ne) | 0 |
| Any other digram | 7 |
| $\mathbf{1 6}$ |  |

## N-Gram Search by Example

- Pattern = "Dauphine" of length $I=8$
- Record = "Universite de Technologie Paris Dauphine"
- $n=2$

- Attempt to match the rightmost 2-gram of pattern against the visited 2-gram in the record
- $\mathrm{AS}(\mathrm{ne})=$ ? AS(si) at offset of "i"


## N-Gram Search by Example

- Pattern = "Dauphine" of length $I=8$
- Record = "Universite de Technologie Paris Dauphine"
- $n=2$

- $67=$ ? 11
- No
- Lookup shift table T at offset $11=$ (AS(si))
- T shows shift of 7 symbols since AS(si) is not in "Dauphine"
- Maximal shift here
- Equal in general to $I-n+1$


## N-Gram Search by Example

- N-Gram Search: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- AS(ne) =? AS( T)
- Mismatch
- What in element AS( T ) in table T ?
- Maximal shift by 7
- Since " T" is nowhere in "Dauphine"


## N-Gram Search by Example

- N-Gram Search: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- Idem
- Mismatch
- Shift by 7
- Again maximal shift since 'lo' not in "Dauphine"


## N-Gram Search by Example

- N-Gram Search: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- Idem
- Mismatch
- Shift by 7
- Maximal shift since 'ar' not in "Dauphine"


## N-Gram Search by Example

- N-Gram Search: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- Compare by signature digrams "ne" and "up"
- Mismatch
- shift by 4 according to $T$
- To align on 'up' in "Dauphine"


## N-Gram Search by Example

- N-Gram Search: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- Match 'ne’ and 'ne’, ‘hi’ and 'hi', ‘up’ against ‘up’, ‘Da' and 'Da'
- Full match


## N-Gram Search by Example

- N-Gram Search: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- Test for false positive : full CAS
- Compare all the matching symbols at the server
- No test if ngram signatures never collide
- e.g., through the method proposed for DNA in the paper


## N-Gram Search by Example

- N-Gram Search: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- Test for false positive : partial CAS
- Compare matching symbols at the server except for AS( $D$ ) in the record
- Match D after decoding at the client
- Remaining $n-1$ leftmost symbols in general
- No test if ngram signatures never collide
- e.g., through the method proposed for DNA in the paper


## BM Search by Example

- Match attempts and shifts compare single symbol at the time

- Compare right-most character
- Mismatch, hence move Dauphine 2 slots to the right where 'i' appears in Dauphine


## BM Search Example

- BM: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:
- Compare right-most character
- Match, hence compare next character
- Mismatch, hence move Dauphine 7 slots to the right since 'e' appears only once in Dauphine


## BM Search Example

- BM: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- Compare 'h' against 'e'
- Mismatch, move pattern three to the right


## BM Search Example

- BM: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- Compare 'l' against 'e'
- No ' l ' in Dauphine, move by 8


## BM Search Example

- BM: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- No 'r' in Dauphine, move by 8


## BM Search Example

- BM: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:
- There is a ' $p$ ' in Dauphine, move by 5


## BM Search Example

- BM: Looking for "Dauphine" in "Universite de Technologie Paris Dauphine:

- Compare 'e' against ' $e$ ', then ' $n$ ' against ' $n$ ', ...
- A match


## Comparison

■ 2-gram search has fewer shifts (6 vs 8)

- The shifts are on average longer
- Even though maximum shift size for 2gram is here only 7 vs. 8 for BM
■ Much larger gain to expect for larger patterns


## N-gram Search in Nutshell



## Performance

- Zero Storage Overhead
$\square$ No indexing
$\square$ Like BM, KMP...
$\square$ Unlike suffix trees and arrays or ngram indexes...
- Search cost is $\mathrm{O}(s)$, $s$ the number of shifts
$\square$ Maximal shift size is $I-n+1$
$\square$ Expected shift size converges towards $f$
- Galois Field size used for CAS calculus


## Performance

- Depends on tuning of $n$
$\square$ Larger $n$ decreases the maximum shift
$\square$ But makes ngrams more discriminative
$\square$ Up to some value of $n$
- depending on the alphabet size, symbol value distribution...
- Our experiments show:
$\square \mathrm{N}=4$ for DNA records
$\square \mathrm{N}=2$ for ASCII \& XML in natural language text


## Analytical Calculus



Expected Shift Size for 4-gram search on DNA

- Random distribution of symbol values


## Experiments

- We compare experimentally performance of N -gram search with BM
- We use mostly partial CAS encoding for:
$\square$ DNA
$\square$ ASCII natural language text
$\square \mathrm{XML}$ code


## Experiments: DNA (homo sap.)



Search Times

## Experiments: DNA (homo sap.)



## Shifts

## Experiments (ASCII nat. lang.)



## Experiments (ASCII nat. lang.)



## Conclusion

- A new algorithm suitable for data stored once and read many times
$\square$ At least as fast as the most used pattern-matching technique (Boyer-Moore);
$\square$ Much faster for small alphabets and/or large patterns;
$\square$ Search without decoding is valuable for P2Pn and Grid environment.
■ Current work on:
$\square$ Approximate string matching
$\square$ Multiple pattern matching
$\square$ Stronger privacy preservation


## Thank You

## for

## Your Attention

