Introduction to Algorithms Lecture 14 String Matching

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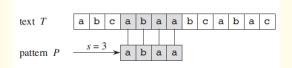
Outline

Introduction

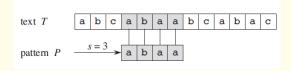
Rabin-Karp Algorithm

3 KMP Algorithm

Given Text $T[1 \dots n]$ and Pattern $P[1 \dots m]$, find all shifts s such that $T[s+1 \dots s+m]=P[1 \dots m]$

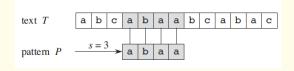


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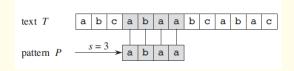
4 Applications in bio-computation (DNA matching), search engine, document processing, ...

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- 2 n and m are large, say $\geq 10^6$
- Any idea of designing a string-matching algorithm?

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- 2 n and m are large, say $\geq 10^6$
- Any idea of designing a string-matching algorithm?
- 4 Our goal is to have time O(n+m)

Preliminaries

- ① Σ : Alphabet with $|\Sigma| = O(1)$ like $\{a, b, ..., z\}$ or $\{0, 1, ..., 9\}$
- ② Σ^n : Strings of length n with alphabet Σ
- $\ \, \mathbf{\mathfrak{I}}^* = \boldsymbol{\Sigma}^0 \cup \boldsymbol{\Sigma}^1 \cup \dots \text{ denotes all strings of alphabet } \boldsymbol{\Sigma}$

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- 4 Prefix $w \sqsubset x$: String w is a prefix of string x, i.e., x = wy for some $y \in \Sigma^*$
- ⑤ Suffix $y \supset x$: String y is a suffix of string x, i.e., x = wy for some $w \in \Sigma^*$

Example: Examples of prefixes and suffixes

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Example: Examples of prefixes and suffixes

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6 Lemma 32.1 in CLRS: If x □ z and y □ z, either x □ y or y □ depends on their length |x| and |y|.

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Recall our problem: Given Text $T[1 \dots n]$ and Pattern $P[1 \dots m]$, find all shifts s such that $T[s+1 \dots s+m] = P[1 \dots m]$.

Basic Idea

When checking T[s+1...s+m] = P[1...m], instead of comparing each element, compare their hash values

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When checking T[s+1...s+m] = P[1...m], instead of comparing each element, compare their hash values

- What kind of hash functions?
- ② How to implement it in time O(n+m)?

① Consider $P[1 \dots m]$ as a huge number in radix-26 $p = P[1] + P[2] \cdot 26 + P[3] \cdot 26^2 + \cdots + P[m] \cdot 26^{m-1}$

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- 4 Consider hash $h: \mathbb{Z} \to \{0, 1, \dots, q-1\}$ like $h[x] = x \mod q$ for $q < 2^{64}$
- ⑤ So $h[P] := p \mod q$ is string P's hash tag.
- 6 How to compute hash tags of $T[1 \dots m], T[2 \dots m+1], \dots, T[n-m+1 \dots n]$?

① Let t_i denote the number T[i+1...i+m], i.e., $t_i := T[i+1] + T[i+2] \cdot 26 + T[i+3] \cdot 26^2 + \cdots + T[i+m] \cdot 26^{m-1}$

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- ② How to compute $t_0 \mod q, \ldots, t_{n-m} \mod q$ efficiently?
- 3 OBS: $h(t_i) = \left(h(t_{i+1}) T[i+1+m] \cdot 26^{m-1}\right) * 26 + T[i+1]$ (mod q) in O(1) time!

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- 4 This gives a linear time algorithm to compute hash tags h(T[1 ... m]), ..., h(T[n-m+1 ... n]) (assume q if not huge)

Analysis

Question

After calculating the hash tags of h[P] and h(T[1 ... m]), ..., h(T[n-m+1 ... n]) in time O(n+m), how to find all shifts s?

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- ① If $t_s \equiv p \mod q$, output "Matching on shift s!"
- Question: Why is it correct?
- 3 Case 1: $t_s = p$ our algorithm will output this shift
- 4 Case 2: $t_s \neq p$ but $h(t_s) = h(p)$ ©
- ⑤ Choose a large prime q in h, say between $[10n^4, 100n^4]$, and consider $\Pr_q[h(t_s) \equiv h(p) \mod q]$ How many primes q are there?

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- ② This happens only if q divides $|t_s p|$.
- 3 How large is $|t_s p|$ and how many prime factors does it have?
- 4 Total number of bad primes is $\leqslant n \cdot \frac{2m}{\log n}$
- None of t_0, \ldots, t_{n-m} gets $h(t_s) = h(p)$ with probability $1 \frac{2nm/\log n}{20n^4/\log n} \geqslant 1 \frac{1}{10n^2}$.

Summary

- ① If we choose a large prime $q \in [10n^4, 100n^4]$, with.prob. $\geqslant 1 \frac{1}{10n^2}$, it outputs all correct shifts s without any mistake.
- Reduce the error probability by taking multiple hash tags

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- 3 Or compare T[i+1...i+m] and P[1...m] when their hash tags are the same O(nm) in worst case but O(n+m) in good cases

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- ① If we choose a large prime $q \in [10n^4, 100n^4]$, with.prob. $\geqslant 1 \frac{1}{10n^2}$, it outputs all correct shifts s without any mistake.
- Reduce the error probability by taking multiple hash tags
- ③ Or compare T[i+1...i+m] and P[1...m] when their hash tags are the same O(nm) in worst case but O(n+m) in good cases
- However, sampling a prime is non-trivial ©
- What if we sample q from [10n⁴, 100n⁴] without the prime constraint?
- Next: KMP algorithms ©

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Switch the idea

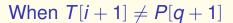
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- ② For each i, consider the longest prefix of P that ends at T[i]

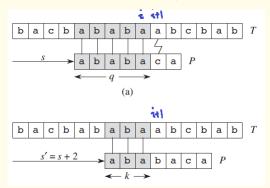
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- ① For each s, find the longest prefix of P starting from T[s+1]
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- 3 Basic idea: Suppose the longest prefix ends at T[i] is P[1 ... q], what if T[i+1] = P[q+1]?

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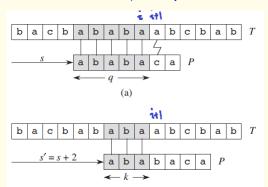
- ① For each s, find the longest prefix of P starting from T[s+1]
- ② For each i, consider the longest prefix of P that ends at T[i]
- 3 Basic idea: Suppose the longest prefix ends at T[i] is P[1 ... q], what if T[i+1] = P[q+1]?
- 4 Key question: What shall we do when $T[i+1] \neq P[q+1]$?





① If we knew the longest prefix of P (again!) that ends at P[q] is P[1 ... k]

When $T[i + 1] \neq P[q + 1]$



- 1 If we knew the longest prefix of P (again!) that ends at P[q] is P[1...k]
- ② We can try comparing T[i+1] with P[k+1]
- ③ Reason: T[i-k+1...i] = P[1...k] since T[i-q+1...i] = P[1...q] and P[1...k] = P[q-k+1...q]

Implementation

Let $\pi[q] = \max \left\{ k : k < q \text{ and } P[1 \dots k] = P[q - k + 1 \dots q] \right\}$ be the longest prefix of P that ends at P[q]

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```
procedure KMP-MATCHER
   q = 0
                                                    // longest prefix at i
   for i = 1, ..., n do
      while q > 0 and P[q + 1] \neq T[i] do
          q = \pi[q]
                                        // Keeping tracing the prefixes
      if P[q + 1] = T[i] then
          q = q + 1
       if q = m then
          output shift at i - m
          q = \pi[q]
                                              // look for the next match
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procedure KMP-MATCHER
   q = 0
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   for i = 1, ..., n do
      while q > 0 and P[q + 1] \neq T[i] do
          a = \pi[a]
                                        // Keeping tracing the prefixes
      if P[q + 1] = T[i] then
          q = q + 1
       if q = m then
          output shift at i - m
          q = \pi[q]
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3 One more question: How to compute π ? — exactly the same problem

```
COMPUTE-PREFIX-FUNCTION (P)
 1 m = P.length
 2 let \pi[1..m] be a new array
 3 \quad \pi[1] = 0
   k = 0
    for q = 2 to m
        while k > 0 and P[k + 1] \neq P[q]
6
            k = \pi[k]
8
     if P[k+1] == P[q]
         k = k + 1
10
    \pi[q] = k
11
    return \pi
```

```
procedure KMP-MATCHER q=0 // longest prefix at i for i=1,\ldots,n do while q>0 and P[q+1]\neq T[i] do q=\pi[q] // Keeping tracing the prefixes if P[q+1]=T[i] then q=q+1 if q=m then output shift at i-m q=\pi[q] // look for the next match
```

Running Time

O(n+m): Apply amortized analysis and control the while loop by a potential function

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- ① $\pi^*[q-1] \subset \{1, 2, ..., q-2\}$ contains all prefixes of P ending at q-1
- **2** Lemma 32.5 in CLRS: If $\pi[1], \ldots, \pi[q-1]$ are correct, $\pi^*[q-1] = \{\pi[q-1], \pi[\pi[q-1]], \ldots, \pi^{(t)}[q-1], \ldots\}$

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- 3 Now consider $\pi[q]$ vs $\pi^*[q-1]$
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- **5** Corollary 32.7 in CLRS: $\pi[q] = \max \left\{ k \in \pi^*[q-1] : P[k+1] = P[q] \right\}$

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Correctness: Basic properties of π

Apply induction on q to show $\pi[q]$ is correct in our ALGO:

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Extending the above proof shows the correctness of KMP

Summary

- Miller-Rabin Algorithm: Easy to implement but relies on random prime numbers
- KMP Algorithm: Tricky to design but fast and reliable
- Automata, data structures for suffixes and prefixes, ...

Questions?

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Break the for-loop of p;