

Algorithms on Strings

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1 String searching

2 Tries

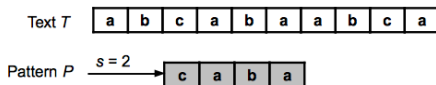
3 Suffix Tries

4 Suffix Trees/Arrays

String searching



- String matching problem: find one or all occurrences of a pattern in a given text
- Applications
 - ▶ information retrieval
 - ▶ Text editors
 - ▶ computational biology (DNA sequences)
- Formal formulation
 - ▶ A text is an array $T[1..n]$ and a pattern is an array $P[1..m]$ ($m \neq n$)
 - ▶ $T[i], P[j] \in$ a finite alphabet Σ (e.g., $\Sigma = \{0, 1\}$ or $\Sigma = \{a, \dots, z\}$)
 - ▶ We say that pattern P **occurs with shift s** in T if $0 \leq s \leq n - m$ and $T[s + 1..s + m] = P[1..m]$



String searching algorithms



- Naive
- Boyer-Moore
- Rabin-Karp
- Knuth-Morris-Pratt (KMP)

Algorithm 1: NaiveSM(P, T)

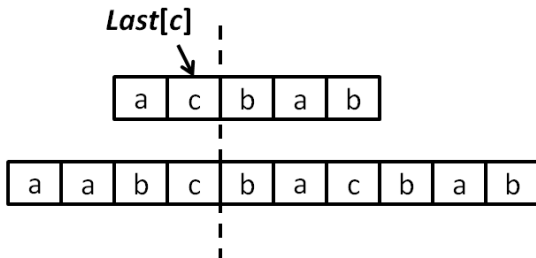
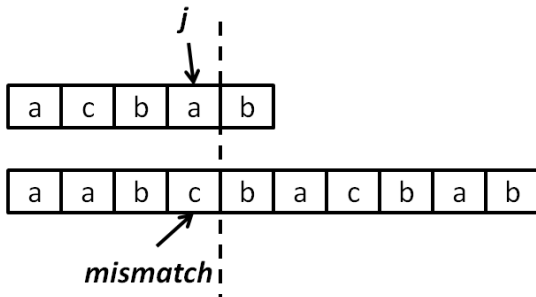
```
foreach  $s = 0..n-m$  do  
   $i \leftarrow 1$ ;  
  while  $i \leq m$  and  $P[i] = T[i + s]$  do  
     $i \leftarrow i + 1$ ;  
  if  $i \geq m$  then  
    Output( $s$ );
```

Boyer-Moore algorithm



- Left to right shift
- Right to left scan
- Use information gained by preprocessing P in order to skip as many alignment as possible
- Bad character shift rule
 - ▶ $last[c]$: the right-most occurrence of c in P
 - ▶ When mismatch: shift P right by $\max\{j - last[c], 1\}$ where j is the position of mismatch character of P

Boyer-Moore algorithm



Boyer-Moore algorithm



```
void computeLast(){
    for(int c = 0; c < 256; c++){
        last[c] = 0;
    }
    for(int i = m; i >= 1; i--){
        if(last[P[i]] == 0)
            last[P[i]] = i;
    }
}

void BoyerMoore(){
    int s = 0;
    while(s <= n-m){
        int j = m;
        while(j > 0 && T[j+s] == P[j]) j--;
        if(j == 0){
            Output(s);
            s = s + 1;
        }else{
            int k = last[T[j+s]];

```


Rabin-Karp algorithm



- Convert the pattern $P[1..m]$ to a number:

$$p = P[1] * d^{m-1} + P[2] * d^{m-2} + \dots + P[m] * d^0$$

where each character $P[i]$ is viewed as a nonnegative integer $< d$, and d is the size of the alphabet

- Using Horner's rule:

$$p = P[m] + d * (P[m-1] + d * (\dots + d * P[1]) + \dots)$$

- Convert $T[s+1..s+m]$ to the integer

$$t_s = T[s+1] * d^{m-1} + \dots + T[s+m]$$

- **Note:** t_{s+1} can easily be computed from t_s as follows:

$$t_{s+1} = (t_s - T[s+1] * d^{m-1}) * d + T[s+m+1]$$

- Drawback: when m is large, then the computation of p and t_s does not take constant time
- Solution: Compute p and t_s modulo a suitable number q
 - ▶ Still problem: $p \equiv t_s \pmod{q}$ does not mean that $p = t_s$, we have to check $P[1..m]$ and $T[s + 1..s + m]$ character by character to see if they are really identical
- Worst-case time is $\mathcal{O}(mn)$ where $P = a^m$ and $T = a^n$

Knuth-Morris-Pratt (KMP) algorithm



- Comparison: from left to right
- Shift: more than one position
- Preprocessing the pattern
 - ▶ Pattern $P[1..m]$
 - ▶ $\pi[q]$ is the length of the longest prefix of $P[1..q]$ which is also the **strictly** suffix of $P[1..q]$

Example

q	1	2	3	4	5	6	7	8	9	10
$P[q]$	a	b	a	b	a	b	a	b	c	a
$\pi[q]$	0	0	1	2	3	4	5	6	0	1

Knuth-Morris-Pratt (KMP) algorithm - preprocessing



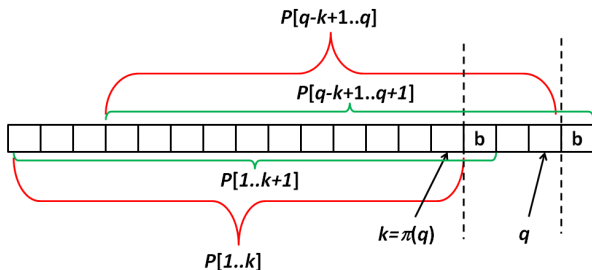
```
void computePI(){
    pi[1] = 0;
    int k = 0;
    for(int q = 2; q <= m; q++){
        while(k > 0 && P[k+1] != P[q])
            k = pi[k];
        if(P[k+1] == P[q])
            k = k + 1;
        pi[q] = k;
    }
}
```

Knuth-Morris-Pratt (KMP) algorithm - preprocessing



Denote $k = \pi[q]$

- If $P[q+1] = P[k+1]$, then $\pi[q+1] = \pi[q] + 1$

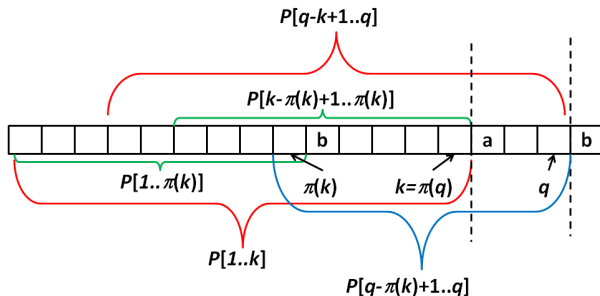


Knuth-Morris-Pratt (KMP) algorithm - preprocessing



Denote $k = \pi[q]$

- if $P[q + 1] \neq P[k + 1]$ and $P[q + 1] = P[\pi[k] + 1] = b$:
 - ▶ $P[1..k] = P[q - k + 1..q] \Rightarrow P[k - \pi[k] + 1..k] = P[q - \pi[k] + 1..q]$
 - ▶ Moreover, $P[k - \pi[k] + 1] = P[1.. \pi[k]]$, so
 $P[1.. \pi[k]] = P[q - \pi[k] + 1..q]$,
 - ▶ Hence $P[1.. \pi[k] + 1] = P[q - \pi[k] + 1..q + 1]$, this means
 $\pi[q + 1] = \pi[k] + 1$

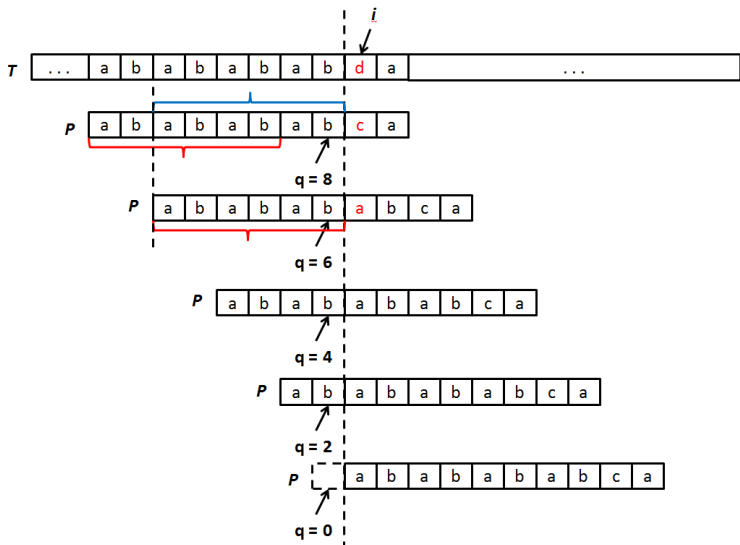


Knuth-Morris-Pratt (KMP) algorithm



```
void kmp(){
    int q = 0;
    for(int i = 1; i <= n; i++){
        while(q > 0 && P[q+1] != T[i]){
            q = pi[q];
        }
        if(P[q+1] == T[i])
            q++;
        if(q == m){
            cout << "match at position " << i-m+1 << endl;
            q = pi[q];
        }
    }
}
```

Knuth-Morris-Pratt (KMP) algorithm



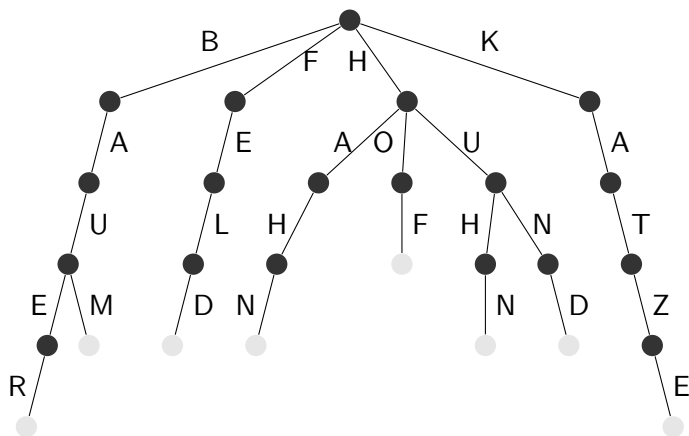
Sets of strings



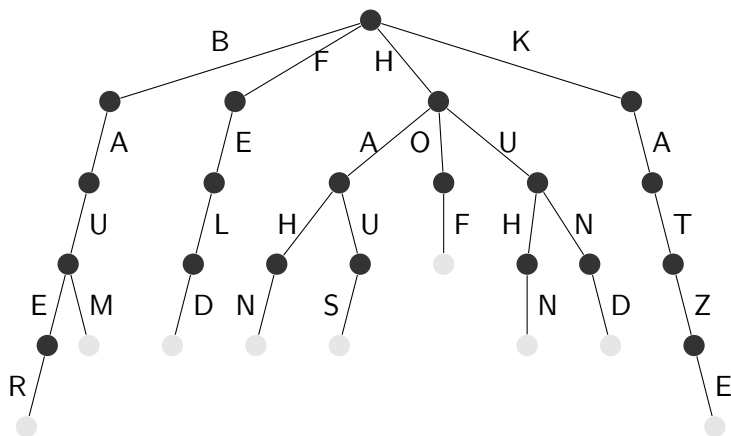
- We often have sets (or maps) of strings
- Insertions and lookups usually guarantee $O(\log n)$ comparisons
- But string comparisons are actually pretty expensive...
- There are other data structures, like tries, which do this in a more clever way

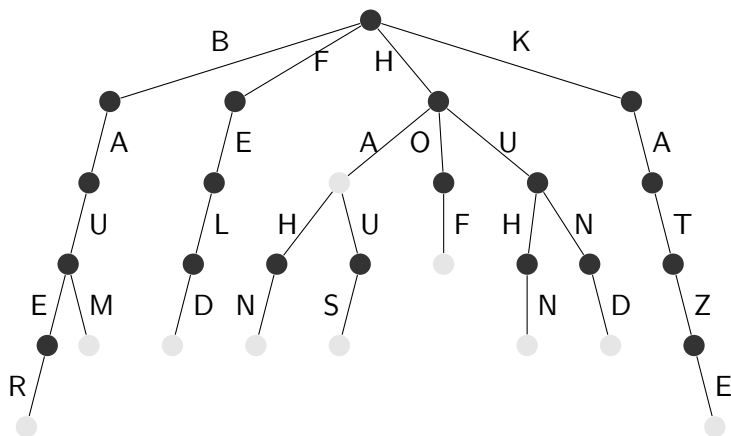
- 1 String searching
- 2 Tries
- 3 Suffix Tries
- 4 Suffix Trees/Arrays

Tries



Tries





```
struct node {  
    node* children[26];  
    bool is_end;  
  
    node() {  
        memset(children, 0, sizeof(children));  
        is_end = false;  
    }  
};
```

```
void insert(node* nd, char *s) {
    if (*s) {
        if (!nd->children[*s - 'a'])
            nd->children[*s - 'a'] = new node();

        insert(nd->children[*s - 'a'], s + 1);
    } else {
        nd->is_end = true;
    }
}
```

Tries



```
bool contains(node* nd, char *s) {  
    if (*s) {  
        if (!nd->children[*s - 'a'])  
            return false;  
  
        return contains(nd->children[*s - 'a'], s + 1);  
    } else {  
        return nd->is_end;  
    }  
}
```


Tries



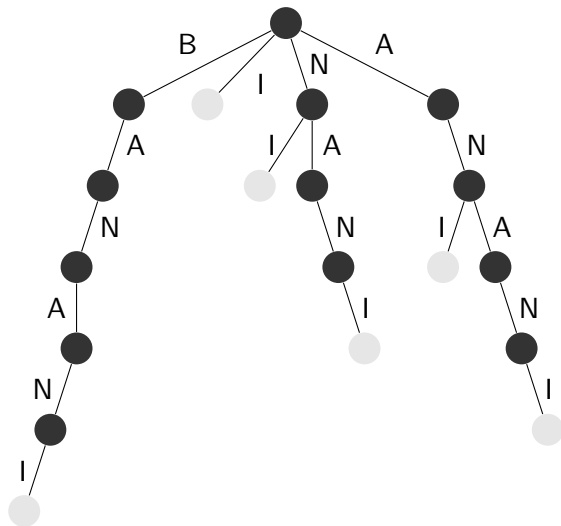
```
node *trie = new node();  
  
insert(trie, "banani");  
  
if (contains(trie, "banani")) {  
    // ...  
}
```

- Time complexity?
- Let k be the length of the string we're inserting/looking for
- Lookup and insertion are both $O(k)$
- Also very space efficient...

- 1 String searching
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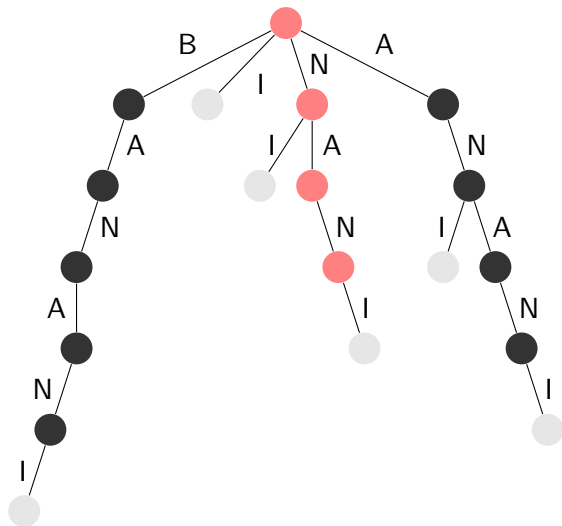
- Say we're dealing with some string S of length n
- Let's insert all suffixes of S into a trie
- $S = \text{banani}$
 - ▶ `insert(trie, "banani");`
 - ▶ `insert(trie, "anani");`
 - ▶ `insert(trie, "nani");`
 - ▶ `insert(trie, "ani");`
 - ▶ `insert(trie, "ni");`
 - ▶ `insert(trie, "i");`

Suffix tries



- There are a lot of cool things we can do with suffix tries
- Example: String matching
- If a string T is a substring in S , then (obviously) it has to start at some suffix of S
- So we can simply look for T in the suffix trie of S , ignoring whether the last node is an end node or not
- This is just $O(m)$...

Suffix tries



- String matching is fast if we have the suffix trie for S
- But what is the time complexity of suffix trie construction?
- There are n suffixes, and it takes $O(n)$ to insert each of them
- So $O(n^2)$, which is pretty slow
- Can we do better?
- There can be up to n^2 nodes in the graph, so this is actually optimal...

1 String searching

2 Tries

3 Suffix Tries

4 Suffix Trees/Arrays

- There exists a compressed version of a suffix trie, called a suffix tree
- It can be constructed in $O(n)$, and has all the features that suffix tries have
- But the $O(n)$ construction algorithm is pretty complex, a big disadvantage for us

Suffix arrays



- A variation of the previous structures
- Can do everything the other structures can do, with a small overhead
- Can be constructed pretty quickly with relatively simple code

Suffix arrays



- Take all the suffixes of S

banani

anani

nani

ani

ni

i

- and sort them

anani

ani

banani

i

nani

ni

Suffix arrays



- We can use this array to do everything that suffix tries can do
- Like string matching

Suffix arrays



- Let's look for nan

anani
ani
banani
i
nani
ni

Suffix arrays



- Let's look for `nan`
- The first letter in the string has to be `n`, so we can binary search for the range of strings starting with `n`

```
anani  
ani  
banani  
i  
nani  
ni
```

Suffix arrays



- Let's look for `nan`
- The first letter in the string has to be `n`, so we can binary search for the range of strings starting with `n`

```
nani  
ni
```


Suffix arrays



- Let's look for `nan`
- The second letter in the string has to be `a`, so we can binary search for the range of strings that have `a` as the second letter

`nani`

`ni`

Suffix arrays



- Let's look for `nan`
- The second letter in the string has to be `a`, so we can binary search for the range of strings that have `a` as the second letter

`nani`

Suffix arrays



- Let's look for `nan`
- The third letter in the string has to be `n`, so we can binary search for the range of strings that have `n` as the third letter

`nani`

Suffix arrays



- Let's look for `nan`
- The third letter in the string has to be `n`, so we can binary search for the range of strings that have `n` as the third letter

`nani`

Suffix arrays



- Let's look for `nan`
- The third letter in the string has to be `n`, so we can binary search for the range of strings that have `n` as the third letter

`nan`
`i`

- If there is at least one string left, we have a match

Suffix arrays



- Time complexity?
- For each letter in T , we do two binary searches on the n suffixes to find the new range
- Time complexity is $O(m \times \log n)$
- A bit slower than doing it with a suffix trie, but still not bad

- But how do we construct a suffix array for a string?
- A simple `sort(suffixes)` is $O(n^2 \log(n))$, because comparing two suffixes is $O(n)$
- And we still have the same problem as with suffix tries, there are almost n^2 characters if we store all suffixes

Suffix arrays



- The second problem is easy to fix
- Just store the indices of the suffixes

```
anani  
ani  
banani  
i  
nani  
ni
```

- becomes

```
1: anani  
3: ani  
0: banani  
5: i  
2: nani  
4: ni
```


- What about the construction?
- In short, we
 - ▶ sort all suffixes by only looking at the first letter
 - ▶ sort all suffixes by only looking at the first 2 letters
 - ▶ sort all suffixes by only looking at the first 4 letters
 - ▶ sort all suffixes by only looking at the first 8 letters
 - ▶ ...
 - ▶ sort all suffixes by only looking at the first 2^i letters
 - ▶ ...
- If we use an $O(n \log n)$ sorting algorithm, this is $O(n \log^2 n)$
- We can also use an $O(n)$ sorting algorithm, since all sorted values are between 0 and n , bringing it down to $O(n \log n)$

Suffix arrays



```
struct suffix_array {
    struct entry {
        pair<int, int> nr;
        int p;

        bool operator <(const entry &other) {
            return nr < other.nr;
        }
    };

    string s;
    int n;
    vector<vector<int> > P;
    vector<entry> L;
    vi idx;

    // constructor
};
```

Suffix arrays



```
suffix_array(string _s) : s(_s), n(s.size()) {
    L = vector<entry>(n);
    P.push_back(vi(n));
    idx = vi(n);

    for (int i = 0; i < n; i++)
        P[0][i] = s[i];

    for (int stp = 1, cnt = 1; (cnt >> 1) < n; stp++, cnt <= 1) {
        P.push_back(vi(n));
        for (int i = 0; i < n; i++) {
            L[i].p = i;
            L[i].nr = make_pair(P[stp - 1][i],
                                i + cnt < n ? P[stp - 1][i + cnt] : -1);
        }
        sort(L.begin(), L.end());
        for (int i = 0; i < n; i++) {
            if (i > 0 && L[i].nr == L[i - 1].nr)
                P[stp][L[i].p] = P[stp][L[i - 1].p];
            else
                P[stp][L[i].p] = i;
        }
    }
    for (int i = 0; i < n; i++)
        idx[P[P.size() - 1][i]] = i;
}
```

Suffix arrays



- There is also one other useful operation on suffix arrays
- Finding the longest common prefix (lcp) of two suffixes of S

```
1: anani
3: ani
0: banani
5: i
2: nani
4: ni
```

- $\text{lcp}(1,3) = 2$
- $\text{lcp}(2,1) = 0$
- This function can be implemented in $O(\log n)$ by using intermediate results from the suffix array construction

Suffix arrays



```
int lcp(int x, int y) {
    int res = 0;
    if (x == y) return n - x;
    for (int k = P.size() - 1; k >= 0 && x < n && y < n; k--)
        if (P[k][x] == P[k][y]) {
            x += 1 << k;
            y += 1 << k;
            res += 1 << k;
        }
    }
    return res;
}
```

Longest common substring

- Given two strings S and T , find their longest common substring
- $S = \text{banani}$
- $T = \text{kanina}$
- Their longest common substring is `ani`

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