



Algorithms Theory

15 – Text search

P.D. Dr. Alexander Souza

Text search

Various scenarios:

Dynamic texts

- Text editors
- Symbol manipulators

Static texts

- Literature databases
- Library systems
- Gene databases
- World Wide Web



Text search

Data type **string**:

- array of character
- file of character
- list of character

Operations (let T, P be of type **string**)

length: $\text{length} ()$

i -th character : $T [i]$

concatenation: $\text{cat} (T, P) \ T.P$

Problem definition

Given:

$$\begin{aligned} \text{text } & t_1 t_2 \dots t_n \in \Sigma^n \\ \text{pattern } & p_1 p_2 \dots p_m \in \Sigma^m \end{aligned}$$

Goal:

Find one or all occurrences of the pattern in the text,
i.e. positions i ($0 \leq i \leq n - m$) such that

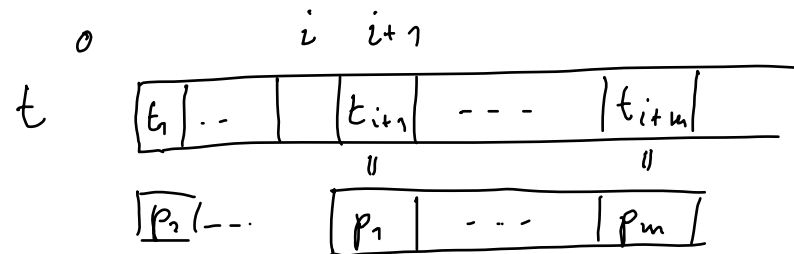
$$p_1 = t_{i+1}$$

$$p_2 = t_{i+2}$$

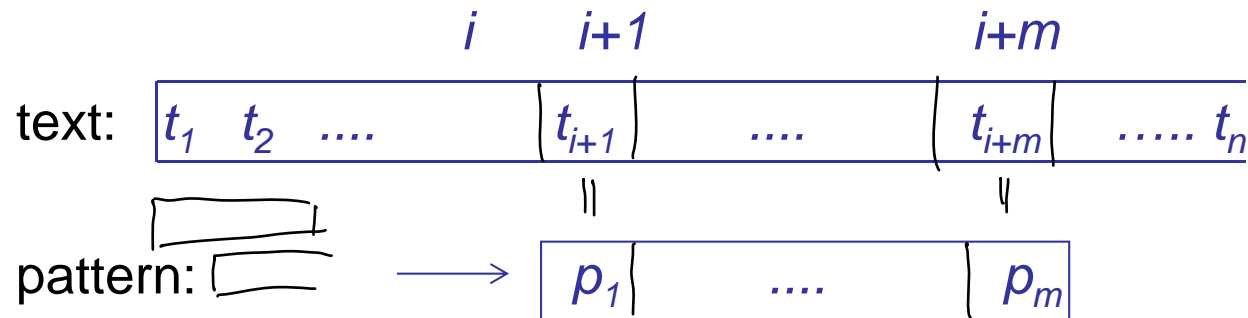
$$\vdots$$

$$p_m = t_{i+m}$$

complete match.



Problem definition



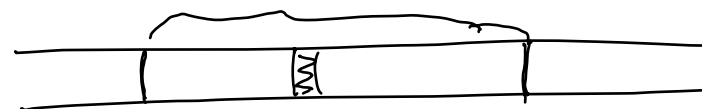
Running time:

1. # possible alignments: $n - m + 1$, # pattern positions: m
 $\rightarrow O(n \cdot m)$

2. At least 1 comparison per m consecutive text positions:

$\rightarrow \Omega(m + n/m)$

reading pattern



≥ 1 comparison for any correct algorithm
 otherwise pattern could be missed.

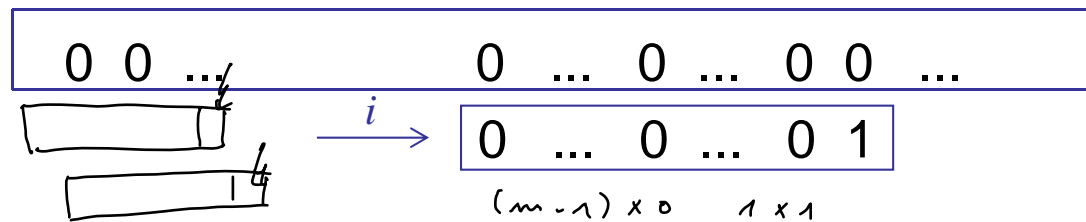
Naive method

For each possible position $0 \leq i \leq n - m$, check at most m character pairs. Whenever a mismatch occurs, shift to the next position.

```
textsearchbf := proc (T :: string, P :: string)
# Input:   text T, pattern P
# Output:  list L of positions i, at which P occurs in T
  n := length (T); m := length (P);
  L := [ ];
  for i from 0 to n - m do
    j := 1;
    while j ≤ m and T[i + j] = P[j ]
      do j := j + 1 od;
    if j = m + 1 then L := [ L [ ], i ] fi;
  od;
  RETURN (L)
end;
```

Naive method

Running time:



all zeros

Worst case: $\Omega(m \cdot n)$

In practice, a mismatch usually occurs very early.

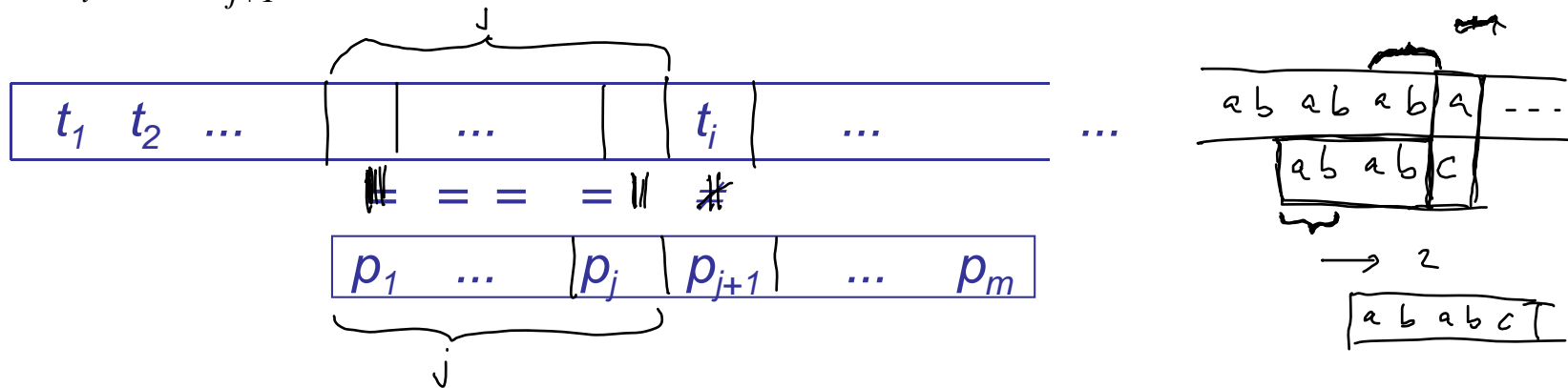
→ running time $\sim cn$ in practice.



The Knuth-Morris-Pratt algorithm (KMP)

We want to find the largest prefix in the pattern which is also a proper suffix.

Let t_i and p_{j+1} be the characters to be compared:



If, for a certain alignment, the first mismatch occurs for characters t_i and p_{j+1} , then:

- the last j characters compared in T equal the first j characters of P
- $t_i \neq p_{j+1}$

Naive: Shift pattern only one position ahead

Idea: Do not necessarily start from ~~scratch~~ scratch, but try to shift the pattern by more than just position. How far?

The Knuth-Morris-Pratt algorithm (KMP)

Idea:

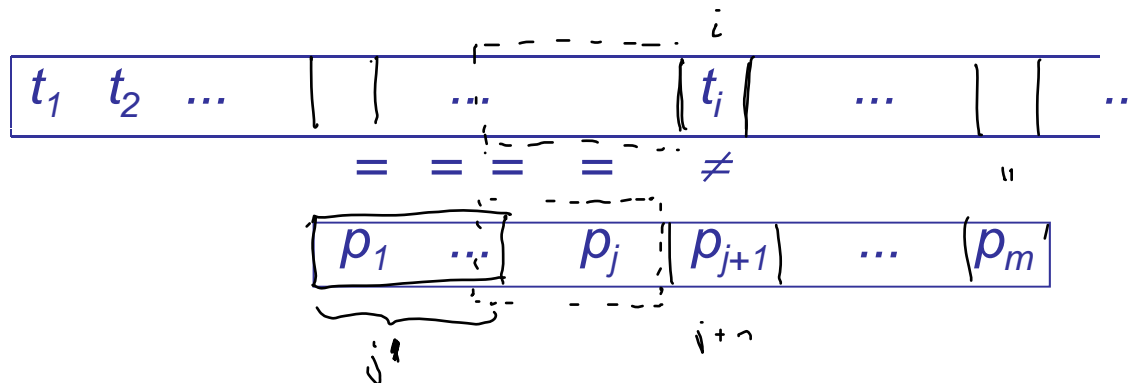
length of the longest prefix of $P_{1\dots j}$ which is also a proper suffix of $P_{1\dots j}$.

array to be constructed

Find $j' = next[j] \leftarrow j$ such that t_i can then be compared to $p_{j'+1}$.

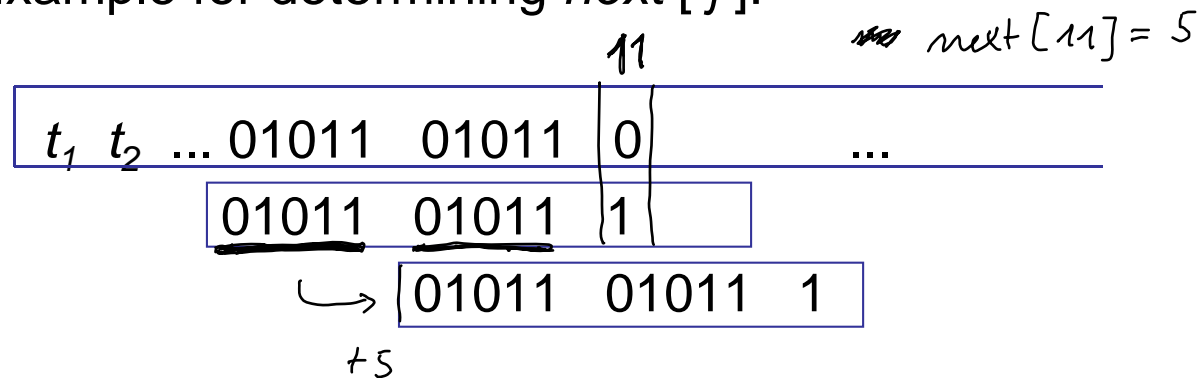
Find greatest $j' < j$ such that $P_{1\dots j'} = P_{j-j'+1\dots j}$.

Find the longest prefix of P that is a proper suffix of $P_{1\dots j}$.



The Knuth-Morris-Pratt algorithm (KMP)

Example for determining $next[j]$:



$next[j] =$ length of the longest prefix of P that is a proper suffix of $P_{1 \dots j}$

The Knuth-Morris-Pratt algorithm (KMP)

\Rightarrow for $P = 0101101011$, $next = [0, 0, 1, 2, 0, 1, 2, 3, 4, 5]$:

1	2	3	4	5	6	7	8	9	10
0	1	0	1	1	0	1	0	1	1
<u>0</u>	<u>1</u>	0	1	1	0	1	0	1	1
<u>0</u>	<u>1</u>	0	1	1	0	1	0	1	1
<u>0</u>	<u>1</u>	0	1	1	0	1	0	1	1
<u>0</u>	<u>1</u>	0	1	1	0	1	0	1	1
<u>0</u>	<u>1</u>	0	1	1	0	1	0	1	1
<u>0</u>	<u>1</u>	0	1	1	0	1	0	1	1
<u>0</u>	<u>1</u>	0	1	1	0	1	0	1	1
<u>0</u>	<u>1</u>	0	1	1	0	1	0	1	1

The Knuth-Morris-Pratt algorithm (KMP)

```

KMP := proc (T :: string, P :: string)
# Input:  text T, pattern P
# Output: list L of positions i at which P occurs in T
  n := length(T);  m := length(P);
  L := [ ];  next := KMPnext(P);
  j := 0;
  for i from 1 to n do
    { while j >= 0 and T[i] <> P[j+1] do j := next[j] od;
      if T[i] = P[j+1] then j := j+1 fi;
      if j = m then L := [ L[], i-m ];
        j := next[j]
      fi;
    }
  od;
  RETURN (L);
end;

```

$next[j] < j$

The Knuth-Morris-Pratt algorithm (KMP)

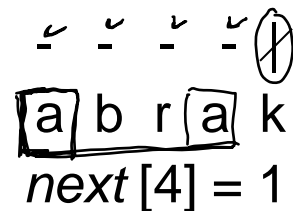
Pattern: abrakadabra, $next = [0, 0, 0, 1, 0, 1, 0, 1, 2, 3, 4]$

a b r a k a d a b r a b r a b a b r a k ... $i = 0$



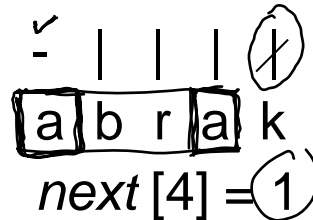
$next[11] = \underline{4}$

a b r a k a d a b r a b r a b a b r a k ... $i = 4$

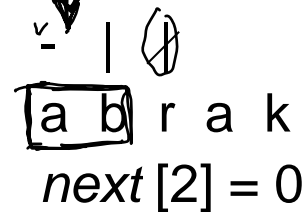


The Knuth-Morris-Pratt algorithm (KMP)

a b r a k a d a b r a b r a b a b r a k ...



a b r a k a d a b r a b r a b a b r a k ...

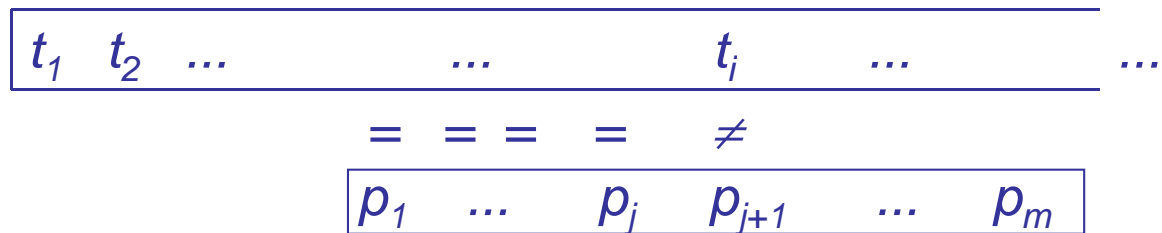


a b r a k a d a b r a b r a b a b r a k ...



The Knuth-Morris-Pratt algorithm (KMP)

Correctness:



When starting the for-loop:

$$P_{1..j} = T_{i-j..i-1} \text{ and } j \neq m$$

if $j = 0$: we are located at the first character of P

if $j \neq 0$: P can be shifted while $j > 0$ and $t_i \neq p_{j+1}$

The Knuth-Morris-Pratt algorithm (KMP)



If $T[i] = P[j+1]$, j and i can be increased (at the end of the loop).

If P has been compared completely ($j = m$), an occurrence of P in T has been found and we can shift to the next position.



The Knuth-Morris-Pratt algorithm (KMP)

Running time:

- the text pointer i is never reset
- text pointer i and pattern pointer j are always incremented together
- always: $next[j] < j$;
 j can be decreased only as many times as it has been increased

If the *next*-array is known, the KMP algorithm runs in $O(n)$ time.

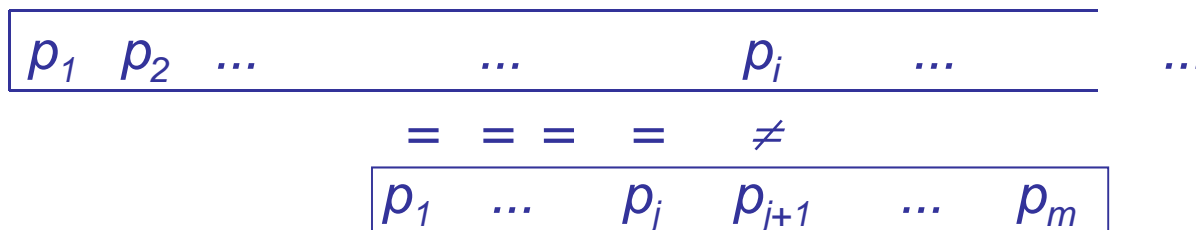
$$\underline{O(n + m)}$$

Computation of the *next*-array

$next[i]$ = length of the longest prefix of P that is a proper suffix of $P_{1\dots i}$

$next[1] = 0$

Let $next[i-1] = j$:



Computation of the *next*-array

Consider two cases:

1) $p_i = p_{j+1} \rightarrow next[i] = j + 1$

2) $p_i \neq p_{j+1} \rightarrow$ replace j by $next[j]$ until $p_i = p_{j+1}$ or $j = 0$
If $p_i = p_{j+1}$, set $next[i] = j + 1$, otherwise $next[i] = 0$.

Computation of the *next*-array

```
KMPnext := proc (P :: string)
# Input:  pattern P
# Output: next-array for P
  m := length (P);
  next := array (1.. m);
  next [1] := 0;
  j := 0;
  for i from 2 to m do
    while j > 0 and P [ i ] <> P [ j+1 ]
      do j := next [ j ] od;
    if P [ i ] = P [ j+1 ] then j := j+1 fi;
    next [ i ] := j
  od;
  RETURN (next);
end;
```



Running time of KMP

The KMP algorithm runs in $O(n + m)$ time.

Can text search be realized even faster?



The Boyer-Moore algorithm (BM)

Idea: For any alignment of the pattern with the text, scan the characters from right to left rather than from left to right.

Example:

```
h e s a i d a b r a k a d a b r a b u t
      |
b u t
```

```
h e s a i d a b r a k a d a b r a b u t
      |
b u t
```


BM: last-occurrence function

For $c \in \Sigma$ and the pattern P let

$\delta [c] :=$ index of the right-most occurrence of c in P

$$\begin{aligned} &= \max \{j \mid p_j = c\} \\ &= \begin{cases} 0 & \text{if } c \notin P \\ j & \text{if } c = p_j \text{ and } c \neq p_k \text{ for } j < k \leq m \end{cases} \end{aligned}$$

What is the cost for computing all δ -values?

Let $|\Sigma| = l$:



BM: last-occurrence function

Let

c = the character causing the mismatch

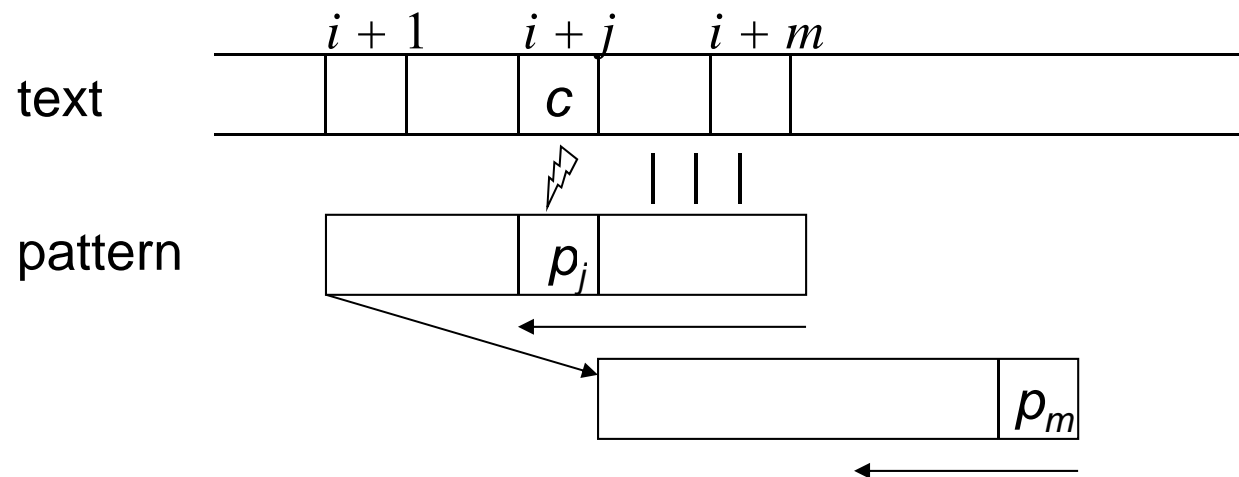
j = the index of the current character in the pattern ($c \neq p_j$)

BM: last-occurrence function

Computation of the pattern shift

Case 1 c does not occur in P ($\delta[c] = 0$)

Shift the pattern j characters to the right.

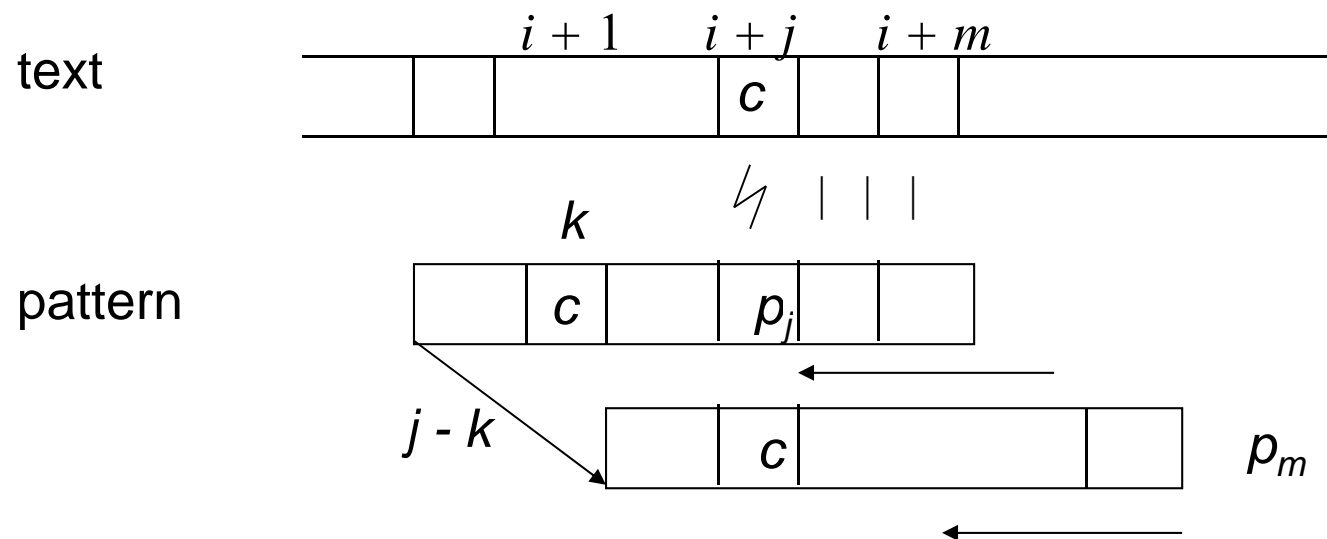


$$\Delta[i] = j$$

BM: last-occurrence function

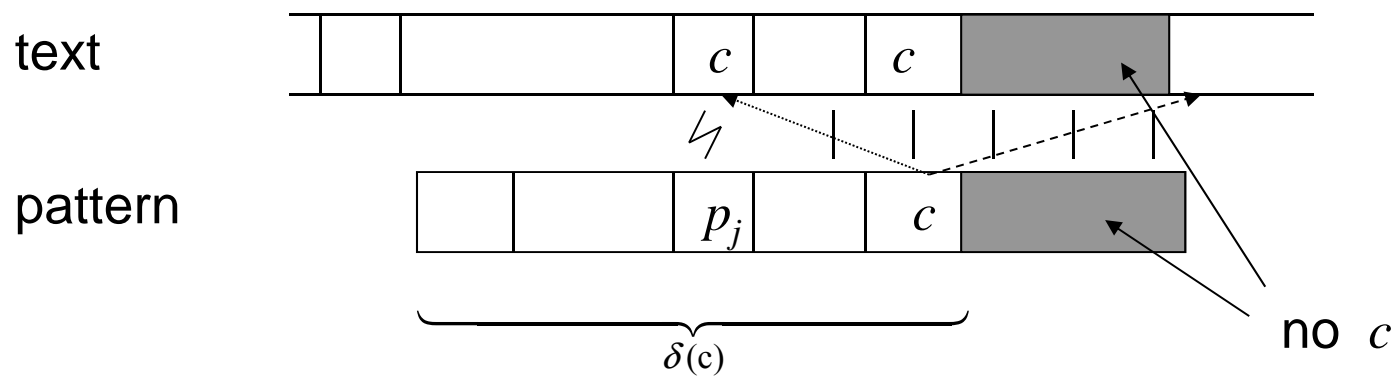
Case 2 c occurs in the pattern ($\delta[c] \neq 0$)

Shift the pattern to the right until the rightmost c in the pattern is aligned with a potential c in the text.



BM: last-occurrence function

Case 2 a: $\delta[c] > j$

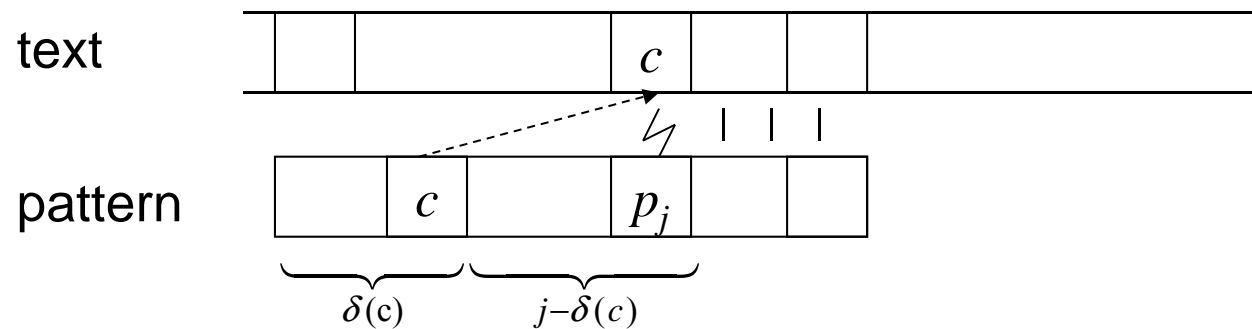


Shift the rightmost c in the pattern to a potential c in the text.

$$\Rightarrow \text{shift by } \Delta[i] = m - \delta[c] + 1$$

BM: last-occurrence function

Case 2 b: $\delta[c] < j$



Shift the rightmost c in the pattern to c in the text.

$$\Rightarrow \text{shift by } \Delta[i] = j - \delta[c]$$

BM: Algorithm (version 1)

Algorithm *BM-search1*

Input: text T , pattern P

Output: all positions of P in T

```
1  $n := \text{length}(T)$ ;  $m := \text{length}(P)$ 
2 compute  $\delta$ 
3  $i := 0$ 
4 while  $i \leq n - m$  do
5      $j := m$ 
6     while  $j > 0$  and  $P[j] = T[i + j]$  do
7          $j := j - 1$ 
8     end while;
```

BM: Algorithm (version 1)

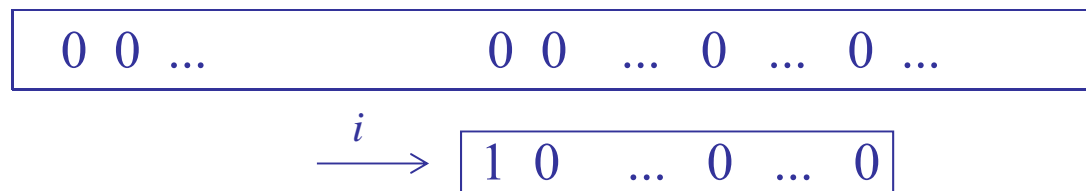
```
8   if  $j = 0$ 
9   then output position  $i$ 
10       $i := i + 1$ 
11   else if  $\delta[ \pi[i + j] ] > j$ 
12      then  $i := i + m + 1 - \delta[ \pi[i + j] ]$ 
13      else  $i := i + j - \delta[ \pi[i + j] ]$ 
14 end while;
```


BM: Algorithm (version 1)

Analysis:

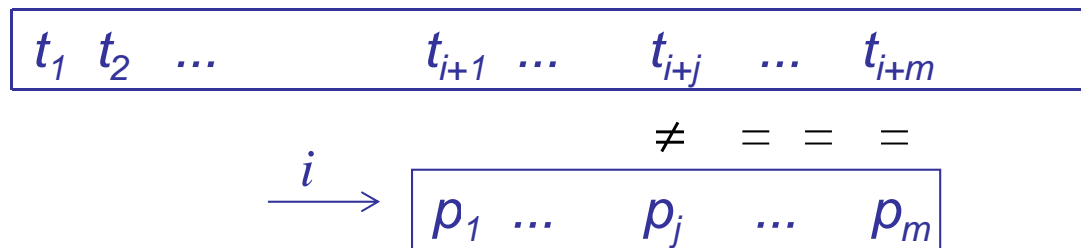
Desired running time: $O(m + n/m)$

Worst-case running time: $\Omega(n m)$



Match heuristic

Use the information collected before a mismatches $p_j \neq t_{i+j}$ occurs.



$gsf[j]$ = position of the end of the next occurrence of the suffix $P_{j+1} \dots m$ from the right that is not preceded by character P_j
 (good suffix function)

Possible shift: $\gamma[j] = m - gsf[j]$

Example of computing gsf

$gsf[j]$ = position of the end of the closest occurrence of the suffix $P_{j+1} \dots m$ from the right that is not preceded by character P_j

pattern: banana

$gsf[j]$	inspected suffix	forbidden character	further occurrence	position
$gsf[5]$	a	n	<u>b</u> an <u>a</u> na	2
$gsf[4]$	na	a	<u>***</u> ba <u>n</u> a <u>na</u>	0
$gsf[3]$	ana	n	ba <u>na</u> na	4
$gsf[2]$	nana	a	ba <u>na</u> na	0
$gsf[1]$	anana	b	<u>ba</u> naana	0

Example of computing gsf

$$\Rightarrow gsf(\text{banana}) = [0,0,0,4,0,2]$$

a b a a b a b a n a n a n a n a

≠ = = =

b a n a n a

b a n a n a

BM: Algorithm (version 2)

Algorithm *BM-search2*

Input: text T , pattern P

Output: shift for all occurrences of P in T

```
1  $n := \text{length}(T)$ ;  $m := \text{length}(P)$ 
2 compute  $\delta$  and  $\gamma$ 
3  $i := 0$ 
4 while  $i \leq n - m$  do
5      $j := m$ 
6     while  $j > 0$  and  $P[j] = T[i + j]$  do
7          $j := j - 1$ 
    end while;
```

BM: Algorithm (version 2)

```
8   if  $j = 0$ 
9     then output position  $i$ 
10       $i := i + \gamma[0]$ 
11   else  $i := i + \max(\gamma[j], j - \delta[T[i+j]])$ 
12 end while;
```