

- The search time of BDM and BOM is $\mathcal{O}(n(\log_{\sigma} m)/m)$, which is optimal on **average**. (BNDM is optimal only when $m \leq w$.)
- MP and KMP are optimal in the **worst case**.
- There are also algorithms that are optimal in **both** cases. They are based on similar techniques, but they will not be described them here.

Karp–Rabin

The Karp–Rabin algorithm uses a **hash function** $H : \Sigma^* \rightarrow [0..q) \subset \mathbb{N}$ for strings. It computes $H(P)$ and $H(T[j..j + m))$ for all $j \in [0..n - m]$.

- If $H(P) \neq H(T[j..j + m))$, then we must have $P \neq T[j..j + m)$.
- If $H(P) = H(T[j..j + m))$, the algorithm compares P and $T[j..j + m)$ in brute force manner. If $P \neq T[j..j + m)$, this is a **collision**.

A good hash function has two important properties:

- Collisions are rare.
- Given $H(a\alpha)$, a and b , where $a, b \in \Sigma$ and $\alpha \in \Sigma^*$, we can quickly compute $H(\alpha b)$. This is called **rolling** or **sliding window** hash function.

The latter property is essential for fast computation of $H(T[j..j + m))$ for all j .

The hash function used by Karp–Rabin is

$$H(c_0c_1c_2 \dots c_{m-1}) = (c_0r^{m-1} + c_1r^{m-2} + \dots + c_{m-2}r + c_{m-1}) \bmod q$$

This is a rolling hash function:

$$H(\alpha) = (H(a\alpha) - ar^{m-1}) \bmod q$$

$$H(\alpha b) = (H(\alpha) \cdot r + b) \bmod q$$

which follows from the rules of [modulo arithmetic](#):

$$(x + y) \bmod q = ((x \bmod q) + (y \bmod q)) \bmod q$$

$$(xy) \bmod q = ((x \bmod q)(y \bmod q)) \bmod q$$

The parameters q and r have to be chosen with some care to ensure that collisions are rare.

- The original choice is $r = \sigma$ and q is a large [prime](#).
- Another possibility is $q = 2^w$, where w is the machine word size, and r is a small prime ($r = 37$ has been suggested). This is faster in practice, because it avoids slow modulo operations.
- The hash function can be [randomized](#) by choosing q or r randomly. Furthermore, we can change q or r whenever a collision happens.

Algorithm 1.17: Karp-Rabin

Input: text $T = T[0 \dots n)$, pattern $P = P[0 \dots m)$

Output: position of the first occurrence of P in T

- (1) Choose q and r ; $s \leftarrow r^{m-1} \bmod q$
- (2) $hp \leftarrow 0$; $ht \leftarrow 0$
- (3) **for** $i \leftarrow 0$ **to** $m - 1$ **do** $hp \leftarrow (hp \cdot r + P[i]) \bmod q$ // $hp = H(P)$
- (4) **for** $j \leftarrow 0$ **to** $m - 1$ **do** $ht \leftarrow (ht \cdot r + T[j]) \bmod q$
- (5) **for** $j \leftarrow 0$ **to** $n - m - 1$ **do**
- (6) **if** $hp = ht$ **then if** $P = T[j \dots j + m)$ **then** return j
- (7) $ht \leftarrow ((ht - T[j] \cdot s) \cdot r + T[j + m]) \bmod q$
- (8) **if** $hp = ht$ **then if** $P = T[j \dots j + m)$ **then** return j
- (9) return n

On an integer alphabet:

- The worst case time complexity is $\mathcal{O}(mn)$.
- The average case time complexity is $\mathcal{O}(m + n)$.

Karp–Rabin is not competitive in practice, but hashing can be a useful technique in other contexts.

2. Approximate String Matching

Often in applications we want to search a text for something that is **similar** to the pattern but not necessarily exactly the same.

To formalize this problem, we have to specify what does “similar” mean. This can be done by defining a **similarity** or a **distance measure**.

A natural and popular distance measure for strings is the **edit distance**, also known as the **Levenshtein distance**.

Edit distance

The **edit distance** $ed(A, B)$ of two strings A and B is the minimum number of edit operations needed to change A into B . The allowed edit operations are:

- S **Substitution** of a single character with another character.
- I **Insertion** of a single character.
- D **Deletion** of a single character.

Example 2.1: Let $A = \text{Lewensteinn}$ and $B = \text{Levenshtein}$. Then $ed(A, B) = 3$.

The set of edit operations can be described with an **edit sequence** or with an **alignment**:

```
NNSNNNINNNND
Lewens-teinn
Levenshtein-
```

In the edit sequence, N means No edit.

There are many variations and extension of the edit distance, for example:

- **Hamming distance** allows only the substitution operation.
- **Damerau–Levenshtein distance** adds an edit operation:
T **Transposition** swaps two adjacent characters.
- With **weighted edit distance**, each operation has a cost or weight, which can be other than one.
- Allow insertions and deletions (indels) of **factors** at a cost that is lower than the sum of character indels.

We will focus on the basic Levenshtein distance.

Computing Edit Distance

Given two strings $A[1..m]$ and $B[1..n]$, define the values d_{ij} with the recurrence:

$$\begin{aligned}d_{00} &= 0, \\d_{i0} &= i, \quad 1 \leq i \leq m, \\d_{0j} &= j, \quad 1 \leq j \leq n, \text{ and} \\d_{ij} &= \min \begin{cases} d_{i-1,j-1} + \delta(A[i], B[j]) \\ d_{i-1,j} + 1 \\ d_{i,j-1} + 1 \end{cases} \quad 1 \leq i \leq m, 1 \leq j \leq n,\end{aligned}$$

where

$$\delta(A[i], B[j]) = \begin{cases} 1 & \text{if } A[i] \neq B[j] \\ 0 & \text{if } A[i] = B[j] \end{cases}$$

Theorem 2.2: $d_{ij} = ed(A[1..i], B[1..j])$ for all $0 \leq i \leq m$, $0 \leq j \leq n$.
In particular, $d_{mn} = ed(A, B)$.

Example 2.3: $A = \text{ballad}, B = \text{handball}$

d		h	a	n	d	b	a	l	l
	0	1	2	3	4	5	6	7	8
b	1	1	2	3	4	4	5	6	7
a	2	2	1	2	3	4	4	5	6
l	3	3	2	2	3	4	5	4	5
l	4	4	3	3	3	4	5	5	4
a	5	5	4	4	4	4	4	5	5
d	6	6	5	5	4	5	5	5	6

$$ed(A, B) = d_{mn} = d_{6,8} = 6.$$

Proof of Theorem 2.2. We use induction with respect to $i + j$. For brevity, write $A_i = A[1..i]$ and $B_j = B[1..j]$.

Basis:

$$\begin{aligned}d_{00} &= 0 = ed(\epsilon, \epsilon) \\d_{i0} &= i = ed(A_i, \epsilon) \quad (i \text{ deletions}) \\d_{0j} &= j = ed(\epsilon, B_j) \quad (j \text{ insertions})\end{aligned}$$

Induction step: We show that the claim holds for d_{ij} , $1 \leq i \leq m$, $1 \leq j \leq n$. By induction assumption, $d_{pq} = ed(A_p, B_q)$ when $p + q < i + j$.

The value $ed(A_i, B_j)$ is based on an **optimal edit sequence**. We have three cases depending on what the last edit operation is:

$$\text{N or S: } ed(A_i, B_j) = ed(A_{i-1}, B_{j-1}) + \delta(A[i], B[j]) = d_{i-1, j-1} + \delta(A[i], B[j]).$$

$$\text{I: } ed(A_i, B_j) = ed(A_i, B_{j-1}) + 1 = d_{i, j-1} + 1.$$

$$\text{D: } ed(A_i, B_j) = ed(A_{i-1}, B_j) + 1 = d_{i-1, j} + 1.$$

Since the edit sequence is optimal, the correct value is the minimum of the three cases, which agrees with the definition of d_{ij} . □

The recurrence gives directly a **dynamic programming** algorithm for computing the edit distance.

Algorithm 2.4: Edit distance

Input: strings $A[1..m]$ and $B[1..n]$

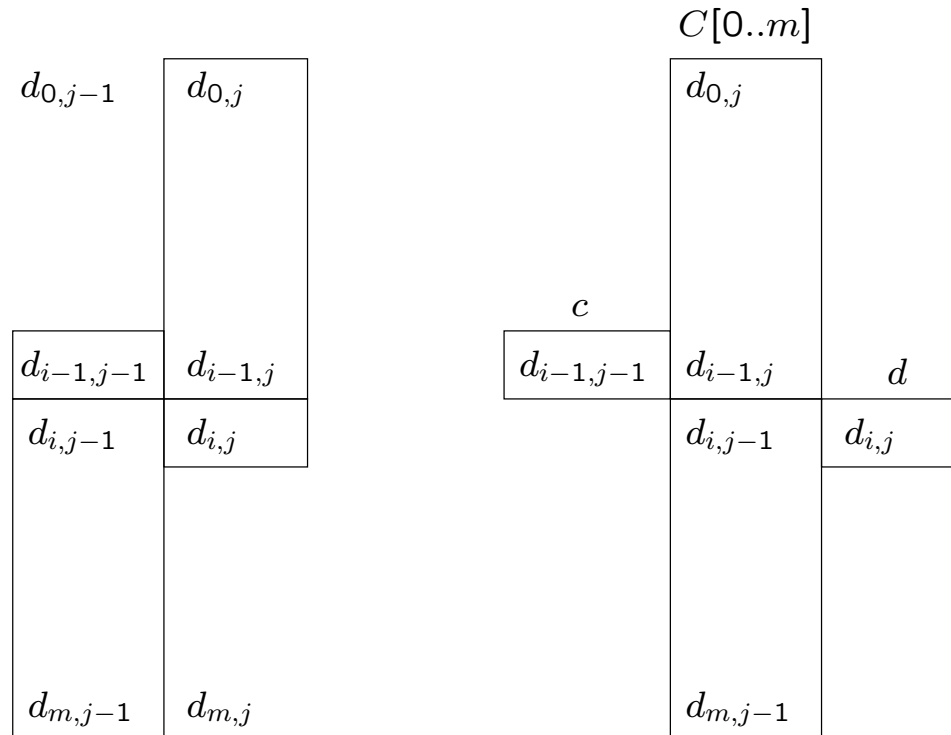
Output: $ed(A, B)$

- (1) for $i \leftarrow 0$ to m do $d_{i0} \leftarrow i$
- (2) for $j \leftarrow 1$ to n do $d_{0j} \leftarrow j$
- (3) for $j \leftarrow 1$ to n do
- (4) for $i \leftarrow 1$ to m do
- (5) $d_{ij} \leftarrow \min\{d_{i-1,j-1} + \delta(A[i], B[j]), d_{i-1,j} + 1, d_{i,j-1} + 1\}$
- (6) return d_{mn}

The time and space complexity is $\mathcal{O}(mn)$.

The space complexity can be reduced by noticing that each column of the matrix (d_{ij}) depends **only on the previous column**. We do not need to store older columns.

A more careful look reveals that, when computing d_{ij} , we only need to store the bottom part of column $j - 1$ and the already computed top part of column j . We store these in an array $C[0..m]$ and variables c and d as shown below:



Algorithm 2.5: Edit distance in $\mathcal{O}(m)$ space

Input: strings $A[1..m]$ and $B[1..n]$

Output: $ed(A, B)$

```
(1) for  $i \leftarrow 0$  to  $m$  do  $C[i] \leftarrow i$ 
(2) for  $j \leftarrow 1$  to  $n$  do
(3)    $c \leftarrow C[0]$ ;  $C[0] \leftarrow j$ 
(4)   for  $i \leftarrow 1$  to  $m$  do
(5)      $d \leftarrow \min\{c + \delta(A[i], B[j]), C[i - 1] + 1, C[i] + 1\}$ 
(6)      $c \leftarrow C[i]$ 
(7)      $C[i] \leftarrow d$ 
(8) return  $C[m]$ 
```

- Note that because $ed(A, B) = ed(B, A)$ (exercise), we can assume that $m \leq n$.

It is also possible to find optimal edit sequences and alignments from the matrix d_{ij} .

An edit graph is a directed graph, where the nodes are the cells of the edit distance matrix, and the edges are as follows:

- If $A[i] = B[j]$ and $d_{ij} = d_{i-1,j-1}$, there is an edge $(i-1, j-1) \rightarrow (i, j)$ labelled with N.
- If $A[i] \neq B[j]$ and $d_{ij} = d_{i-1,j-1} + 1$, there is an edge $(i-1, j-1) \rightarrow (i, j)$ labelled with S.
- If $d_{ij} = d_{i,j-1} + 1$, there is an edge $(i, j-1) \rightarrow (i, j)$ labelled with I.
- If $d_{ij} = d_{i-1,j} + 1$, there is an edge $(i-1, j) \rightarrow (i, j)$ labelled with D.

Any path from $(0,0)$ to (m,n) is labelled with an optimal edit sequence.

Example 2.6: $A = \text{ballad}, B = \text{handball}$

d		h	a	n	d	b	a	l	l
	0	⇒ 1	⇒ 2	⇒ 3	⇒ 4	→ 5	→ 6	→ 7	→ 8
b	↓	↘	↘	↘	↘	↘			
	1	1	→ 2	→ 3	→ 4	4	→ 5	→ 6	→ 7
a	↓	↘	↓	↘			↘		
	2	2	1	⇒ 2	→ 3	→ 4	4	→ 5	→ 6
l	↓	↘	↓	↘	↘	↘	↘	↓	↘
	3	3	2	2	⇒ 3	→ 4	→ 5	4	→ 5
l	↓	↘	↓	↘	↘	↘	↘	↘	↘
	4	4	3	3	3	⇒ 4	→ 5	5	4
a	↓	↘	↓	↘	↓	↘	↘	↘	↘
	5	5	4	4	4	4	4	⇒ 5	5
d	↓	↘	↓	↘	↓	↘	↓	↘	↘
	6	6	5	5	4	→ 5	5	5	⇒ 6

There are 7 paths from (0,0) to (6,8) corresponding to, for example, the following edit sequences and alignments:

IIIIINNND	SNISSNIS	SNSSINSI
----ballad	ba-lla-d	ball-ad-
handball--	handball	handball