

Algoritmi per la Bioinformatica

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String Distance Measures

Similarity vs. distance

Two ways of measuring the same thing:

1. How **similar** are two strings?
 2. How **different** are two strings?
1. **Similarity**: the **higher** the value, the closer the two strings.
 2. **Distance**: the **lower** the value, the closer the two strings.

Similarity vs. distance

Example

s = TATTACTATC
t = CATTAGTATC

- number of equal positions: $|\{i : s_i = t_i\}| = 8$ (out of 10)
80% **similarity** ($s = t$ if 100%, i.e. if **high**)
- number of different positions: $|\{i : s_i \neq t_i\}| = 2$ (out of 10)
Hamming **distance** 2 ($s = t$ if 0, i.e. if **low**)

(Note that both are defined only if $|s| = |t|$.)

Alignment score and edit distance

Edit operations

- **substitution**: a becomes b , where $a \neq b$
- **deletion**: delete character a
- **insertion**: insert character a

Often one views alignments in this way:

ACCT	ACCT--	-ACCT
CACT	--CACT	CA-CT
2 substitutions	2 deletions, 1 substitution, 2 insertions	1 insertion, 1 deletion

The edit distance

Edit distance, also called **Levenshtein distance**, or unit-cost edit distance (Levenshtein, 1965)

Definition

The edit distance $d(s, t)$ is the **minimum** number of edit operations needed to transform s into t .

Example

s = TACAT, t = TGATAT

- TACAT $\xrightarrow{\text{subst}}$ GACAT $\xrightarrow{\text{del}}$ GAAT $\xrightarrow{\text{ins}}$ TGAAT $\xrightarrow{\text{ins}}$ TGATAT 4 edit op's

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- TACAT $\xrightarrow{\text{ins}}$ TGACAT $\xrightarrow{\text{subst}}$ TGAGAT $\xrightarrow{\text{subst}}$ TGATAT 3 edit op's

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Example

$s = \text{TACAT}$, $t = \text{TGATAT}$

- $\text{TACAT} \xrightarrow{\text{subst}} \text{GACAT} \xrightarrow{\text{del}} \text{GAAT} \xrightarrow{\text{ins}} \text{TGAAT} \xrightarrow{\text{ins}} \text{TGATAT}$ 4 edit op's
- $\text{TACAT} \xrightarrow{\text{ins}} \text{TGACAT} \xrightarrow{\text{subst}} \text{TGAGAT} \xrightarrow{\text{subst}} \text{TGATAT}$ 3 edit op's
- $\text{TACAT} \xrightarrow{\text{ins}} \text{TGACAT} \xrightarrow{\text{subst}} \text{TGATAT}$ 2 edit op's

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Alignments vs. edit operations

Not every series of operations corresponds to an alignment:

- $\text{TACAT} \xrightarrow{\text{subst}} \text{GACAT} \xrightarrow{\text{del}} \text{GAAT} \xrightarrow{\text{ins}} \text{TGAAT} \xrightarrow{\text{ins}} \text{TGATAT}$
- $\text{TACAT} \xrightarrow{\text{ins}} \text{TGACAT} \xrightarrow{\text{subst}} \text{TGAGAT} \xrightarrow{\text{subst}} \text{TGATAT}$
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-TAC-AT
TGA-TAT
- $\text{TACAT} \xrightarrow{\text{ins}} \text{TGACAT} \xrightarrow{\text{subst}} \text{TGAGAT} \xrightarrow{\text{subst}} \text{TGATAT}$
???
- $\text{TACAT} \xrightarrow{\text{ins}} \text{TGACAT} \xrightarrow{\text{subst}} \text{TGATAT}$
T-ACAT
TGATAT

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Alignments vs. edit operations

But every alignment corresponds to a series of operations:

- match \mapsto do nothing
- mismatch \mapsto substitution
- gap below \mapsto deletion
- gap on top \mapsto insertion

Example

T-ACAT-
TGAT-AT

$\text{TACAT} \xrightarrow{\text{ins}} \text{TGACAT} \xrightarrow{\text{subst}} \text{TGATAT} \xrightarrow{\text{del}} \text{TGATT} \xrightarrow{\text{subst}} \text{TGATA} \xrightarrow{\text{ins}} \text{TGATAT}$

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Alignments vs. edit operations

Take the following scoring function: $\text{match} = 0$, $\text{mismatch} = -1$, $\text{gap} = -1$. If alignment \mathcal{A} corresponds to the series of operations \mathcal{S} , then:

$$\text{score}(\mathcal{A}) = -|\mathcal{S}|$$

where $|\mathcal{S}| = \text{no. of operations in } \mathcal{S}$.

Example

- $\text{TACAT} \xrightarrow{\text{subst}} \text{GACAT} \xrightarrow{\text{del}} \text{GAAT} \xrightarrow{\text{ins}} \text{TGAAT} \xrightarrow{\text{ins}} \text{TGATAT}$
-TAC-AT
TGA-TAT
- $\text{TACAT} \xrightarrow{\text{ins}} \text{TGACAT} \xrightarrow{\text{subst}} \text{TGATAT}$
T-ACAT
TGATAT

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Minimum length (shortest) series of edit operations

We are looking for a series of operations of **minimum length**:

$\text{dist}(s, t) = \min\{|\mathcal{S}| : \mathcal{S} \text{ is a series of operations transforming } s \text{ into } t\}$

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Exercises on edit distance

Exercises

- If t is a substring of s , then what is $\text{dist}(s, t)$?
- What is $\text{dist}(s, \epsilon)$?
- If we can transform s into t by using only deletions, then what can we say about s and t ?
- If we can transform s into t by using only substitutions, then what can we say about s and t ?

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What is a distance?

A **distance function** (metric) on a set X is a function $d : X \times X \rightarrow \mathbb{R}$ s.t. for all $x, y, z \in X$:

1. $d(x, y) \geq 0$, and $d(x, y) = 0 \Leftrightarrow x = y$ (positive definite)
2. $d(x, y) = d(y, x)$ (symmetric)
3. $d(x, y) \leq d(x, z) + d(z, y)$ (triangle inequality)

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Examples

- Euclidean distance on \mathbb{R}^2 : $d(x, y) = \sqrt{(x_1 - y_1)^2 + (x_2 - y_2)^2}$
- Manhattan distance on \mathbb{R}^2 : $d(x, y) = |x_1 - y_1| + |x_2 - y_2|$
- Hamming distance on Σ^n : $d_H(s, t) = \{i : s_i \neq t_i\}$.

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The edit distance is a distance

The edit distance is a metric (distance function):

Let $s, t, u \in \Sigma^*$ (strings over Σ):

1. $\text{dist}(s, t) \geq 0$: to transform s into t , we need 0 or more edit op's. Also, we can transform s into t with 0 edit op's if and only if $s = t$.
2. Since every edit operation can be inverted, we get $\text{dist}(s, t) = \text{dist}(t, s)$.
3. (by contradiction) Assume that $\text{dist}(s, u) + \text{dist}(u, t) < \text{dist}(s, t)$, and S transforms s into u in $\text{dist}(s, u)$ steps, and S' transforms u into t in $\text{dist}(u, t)$ steps. Then the series of op's $S' \circ S$ (first S , then S') transforms s into t , but is shorter than $\text{dist}(s, t)$, a contradiction to the definition of dist .

(Exercise: Show that the Hamming distance is a metric.)

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Computing the edit distance

Note first that we can assume that edit operations happen left-to-right. As for computing an optimal alignment, we look at what happens to the last characters. Transforming s into t can be done in one of 3 ways:

1. transform $s_1 \dots s_{n-1}$ into t and then delete last character of s
2. if $s_n = t_m$: transform $s_1 \dots s_{n-1}$ into $t_1 \dots t_{m-1}$
if $s_n \neq t_m$: transform $s_1 \dots s_{n-1}$ into $t_1 \dots t_{m-1}$ and substitute s_n with t_m
3. transform s into $t_1 \dots t_{m-1}$ and insert t_m

So again we can use Dynamic Programming!

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Computing the edit distance

We will need a DP-table (matrix) E of size $(n+1) \times (m+1)$ (where $n = |s|$ and $m = |t|$).

Definition: $E(i, j) = \text{dist}(s_1 \dots s_i, t_1 \dots t_j)$

Computation of $E(i, j)$:

- Fill in first row and column: $E(0, j) = j$ and $E(i, 0) = i$
- for $i, j > 0$: now $E(i, j)$ is the **minimum** of 3 entries plus 1 or plus 0, depending (on what?)
- return entry on bottom right $E(n, m)$
- backtrack for shortest series of edit operations

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Algorithm for computing the edit distance

Algorithm DP algorithm for edit distance

Input: strings s, t , with $|s| = n, |t| = m$

Output: value $dist(s, t)$

1. **for** $j = 0$ to m **do** $E(0, j) \leftarrow j$;
2. **for** $i = 1$ to n **do** $E(i, 0) \leftarrow i$;
3. **for** $i = 1$ to n **do**

$$E(i, j) \leftarrow \min \begin{cases} E(i-1, j) + 1 \\ E(i-1, j-1) & \text{if } s_i = t_j \\ E(i-1, j-1) + 1 & \text{if } s_i \neq t_j \\ E(i, j-1) + 1 \end{cases}$$

5. **return** $E(n, m)$;

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Analysis

- **Space:** $O(nm)$ for the DP-table
- **Time:**
 - computing $dist(s, t)$: $3nm + n + m + 1 \in O(nm)$ (resp. $O(n^2)$ if $n = m$)
 - finding an optimal series of edit op's: $O(n + m)$ (resp. $O(n)$ if $n = m$)

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Again alignment vs. edit distance

$sim(s, t)$ vs. $dist(s, t)$

Recall the scoring function from before:

$match = 0, mismatch = -1, gap = -1$. Then we have:

$$sim(s, t) = -dist(s, t)$$

(This seems obvious but it actually needs to be proved. Formal proof see Setubal & Meidanis book, Sec. 3.6.1.)

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General cost functions

General cost edit distance: different edit operations can have different cost (but some conditions must hold, e.g. $cost(insert) = cost(delete)$, why?). Also computable with same algorithm in same time and space.

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LCS distance

Given two strings s and t ,

$$LCS(s, t) = \max\{|u| : u \text{ is a subsequence of } s \text{ and } t\}$$

is the length of a longest common subsequence of s and t .

Example

Let $s = \text{TACAT}$ and $t = \text{TGATAT}$

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Example

Let $s = \text{TACAT}$ and $t = \text{TGATAT}$, then we have $LCS(s, t) = 4$.
 $s = \text{TACAT}$, $t = \text{TGATAT}$

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Let $s = \text{TACAT}$ and $t = \text{TGATAT}$, then we have $LCS(s, t) = 4$.
 $s = \text{TACAT}$, $t = \text{TGATAT}$

LCS-distance

$$d_{LCS}(s, t) = |s| + |t| - 2LCS(s, t)$$

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LCS distance

N.B.

There may be more than one longest common subsequence, but the *length* $LCS(s, t)$ is unique! E.g. $s' = \text{TAACAT}$, $t' = \text{ATCTA}$, then $LCS(s', t') = 3$, and ACA , TCA , TCT , ACT are all longest common subsequences.

Example

In the examples above, we have $d_{LCS}(s, t) = 5 + 6 - 2 \cdot 4 = 3$, and $d_{LCS}(s', t') = 6 + 5 - 2 \cdot 3 = 5$.

Exercise (*)

(1) Prove or disprove that this is a metric. (2) Find a DP-algorithm that computes $LCS(s, t)$.

(*) means: for particularly motivated students

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Summary: Similarity and distance

Similarity measures for strings

- $sim(s, t)$ - score of an optimal alignment of s, t
- percent similarity (only for equal length strings!)

Distance measures for strings

- edit distance (Levenshtein distance) - minimum no. of edit operations to transform s into t
- Hamming distance (only for equal length strings!)
- LCS distance
- (q -gram distance)

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Summary: Similarity and distance

- two ways of expressing the same thing (similarity vs. distance)
- **similarity**: the **higher** the value, the more similar the strings
- **distance**: the **lower** the value, the more similar the strings
- optimal alignment \cong minimum length edit transformation
- both computable in quadratic time and quadratic space

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