

Plán semestra

Dnes: *editačná vzdialenosť, najdlhšia spoločná podpostupnosť*

Do pondelka: výber článku na prezentáciu

Streda 17.4.: *zlepšenia výpočtu editačnej vzdialenosti*

Štvrtok 18.4.: prednáška nebude

Streda 24.4.: *približné výskyty vzorky, lokálne podobnosti, bioinformatika*

Štvrtok 25.4.: *zostavovanie DNA sekvencií, najkratšie spoločné nadslovo*

Streda 1.5.: sviatok

Štvrtok 2.5.: *viacnásobné zarovnanie, opakujúce sa sekvenčné motívy*

Streda 8.5.: sviatok

Štvrtok 9.5.: prezentácie

Streda 15.5.: prezentácie

Štvrtok 16.5.: prezentácie

New topic: Approximate occurrences, similar strings

Differences in texts occurs due to typos, experimental errors, transmission errors, deliberately introduced, . . .

So far: Pattern matching with Hamming distance $\leq k$ in $O(nk)$

Today: edit distance between two strings

Edit distance, Levenshtein distance (editačná vzdialenosť')

Edit operations: ($u, v \in \Sigma^*$, $a, b \in \Sigma$)

- insertion (inzercia) $uv \rightarrow uav$
- deletion (delécia) $uav \rightarrow uv$
- substitution (substitúcia) $uav \rightarrow ubv$

Edit distance $d_E(S, T) =$

shortest sequence of edit operations that changes S to T

Example:

$S = \text{ema ma mamu}$, $T = \text{mama sa ma}$, $d_E(S, T) = 5$

ema_ma_mamu (delete e) ma_ma_mamu (delete space)

mama_mamu (substitute m->s) mama_samu (insert space)

mama_sa_mu (substitute u-a) mama_sa_ma

Edit distance as an alignment

Sequence alignment (zarovnanie):

insert gaps (—) to S and T to get matrix with 2 rows

— column with a gap (insertion or deletion): cost 1

— column with two different symbols (substitution): cost 1

— column with equal symbols: cost 0

Example:

ema_ma_ma-mu

-ma-ma_sa_ma

100100010101

Problem: compute $d_E(S, T)$ for two input strings S and T
(note $d_H(S, T)$ trivially in $O(n)$ time)

Dynamic programming for $d_E(S, T)$

Let $m = |S|$, $n = |T|$, indexing from 1: $S[1..m]$, $T[1..n]$

Let $A[i, j] = d_E(S[1..i], T[1..j])$

Compute $A[i, j]$ for $0 \leq i \leq m$, $0 \leq j \leq n$

Example:

12345678901

ema_ma_mamu

mama_sa_ma

$A[3, 4] = 2$

Dynamic programming for $d_E(S, T)$

Let $m = |S|$, $n = |T|$, indexing from 1: $S[1..m]$, $T[1..n]$

Let $A[i, j] = d_E(S[1..i], T[1..j])$

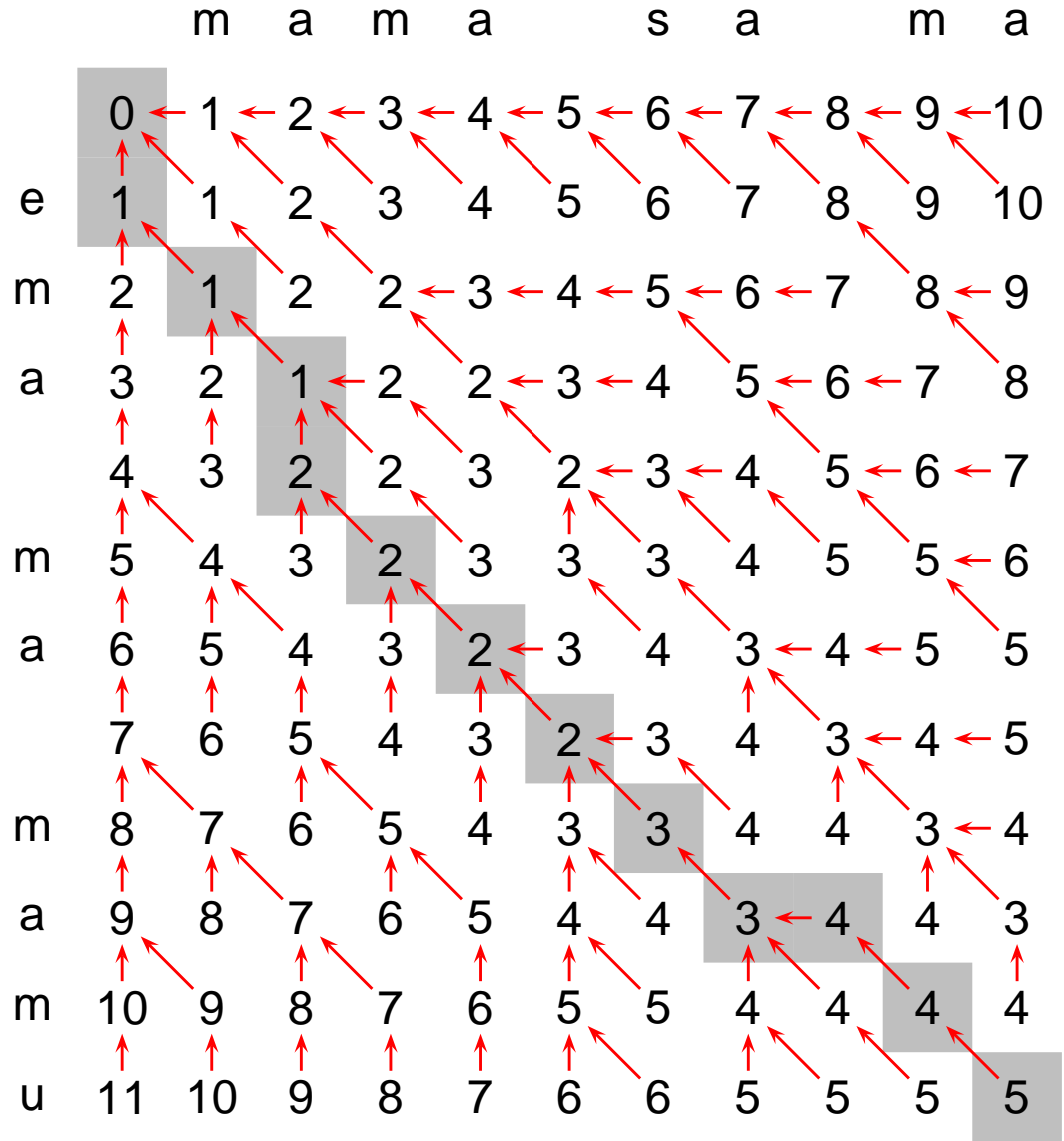
Compute $A[i, j]$ for $0 \leq i \leq m$, $0 \leq j \leq n$

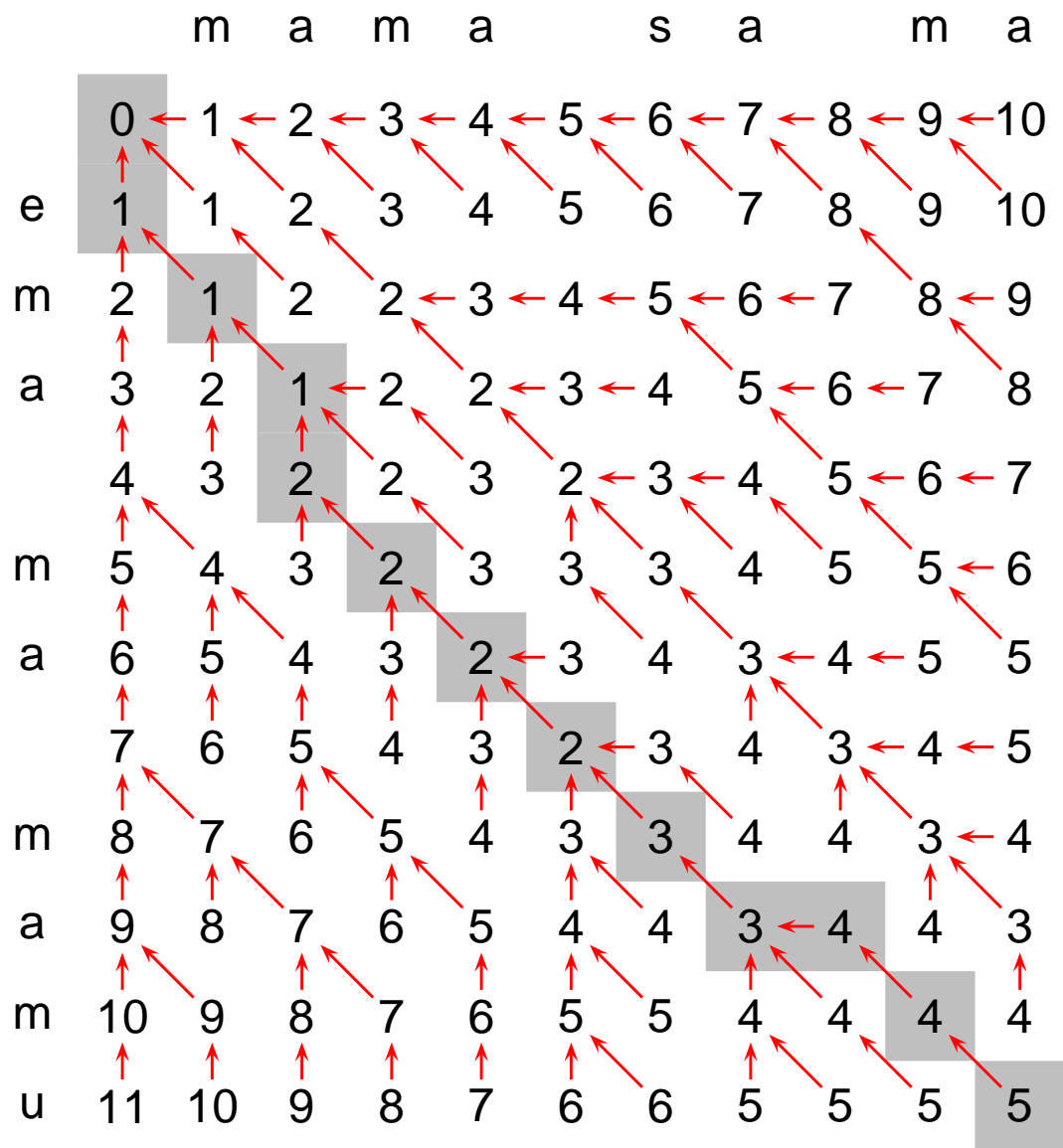
$$A[i, j] = \min \begin{cases} A[i-1, j-1] + c(S[i], T[j]) \\ A[i-1, j] + 1 \\ A[i, j-1] + 1 \end{cases}$$

$$c(a, b) = \begin{cases} 0 & \text{if } a = b \\ 1 & \text{if } a \neq b \end{cases}$$

		m	a	m	a		s	a		m	a
	0	1	2	3	4	5	6	7	8	9	10
e	1	1	2	3	4	5	6	7	8	9	10
m	2	1	2	2	3	4	5	6	7	8	9
a	3	2	1	2	2	3	4	5	6	7	8
	4	3	2	2	3	2	3	4	5	6	7
m	5	4	3	2	3	3	3	4	5		
a											
m											
a											
m											
u											

$$A[i, j] = \min \begin{cases} A[i-1, j-1] \\ \quad + c(S[i], T[j]) \\ A[i-1, j] + 1 \\ A[i, j-1] + 1 \end{cases}$$





Alignment:

ema_ma_ma-mu

-ma-ma_sa_ma

Edit distance is a distance function (metrika)

$$d_E(S, T) \geq 0$$

$$d_E(S, T) = 0 \iff S = T$$

$$d_E(S, T) = d_E(T, S) \text{ (symmetry)}$$

$$d_E(S, T) \leq d_E(S, X) + d_E(X, T) \text{ (triangle inequality)}$$

Generalized edit distance

(zovšeobecnená editačná vzdialenosť)

Table of weights $w(a, b)$ for $a, b \in \Sigma \cup \{-\}$

For $a, b \in \Sigma$:

$w(a, -)$ cost of deletion, $w(-, b)$ cost of insertion

$w(a, b)$ cost of substitution from a to b , or cost of identity if $a = b$

Dynamic programming to find the alignment with lowest cost

– for some strange weights not the lowest sequence of operations

– not always a distance function

$$A[i, j] = \min \begin{cases} A[i-1, j-1] + w(S[i], T[j]) \\ A[i-1, j] + w(S[i], -) \\ A[i, j-1] + w(-, T[j]) \end{cases}$$

Longest common subsequence lcs

Najdlhšia spoločná podpostupnosť

Def: subsequence of a sequence a_1, \dots, a_n is sequence $a_{i_1}, a_{i_2}, \dots, a_{i_k}$ such that $1 \leq i_1 < i_2 < \dots < i_k \leq n$

$lcs(S, T)$ = length of the longest sequence which is a common subsequence of both S and T

Example:

$S = \text{ema ma mamu}, T = \text{mama sa ma}, lcs(S, T) = 7$

Also alignment (disallow substitutions):

```
ema_ma_---mamu
-ma-ma_sama--
. * * . * * * . . . * * . .
```

How to find using DP for generalized edit distance?

Program diff

Compare two files line by line, e.g. two versions of a source code.

Find (approximation of) $lcs(S,T)$ where symbols are lines of the files.

```
1522a1540
```

```
>     my $last_line = undef;
```

```
1525,1526c1543,1544
```

```
<     foreach my $gtf_line (@$transcript) {
```

```
<         # printf STDERR Dumper($gtf_line);
```

```
---
```

```
>     for(my $line_num = 0; $line_num<@$transcript; $line_num++)
```

```
>         my $gtf_line = $transcript->[$line_num];
```

Hunt, Szymanski 1977

$Z[i]$ = list of j s.t. $T[j] = S[i]$ in decreasing order

$Z = Z[1]Z[2] \dots Z[m]$

Example:

12345678901

S: ema_ma_mamu

T: mama_sa_ma

$Z[1]_e = \emptyset$

$Z[2]_m = 9, 3, 1$

$Z[3]_a = 10, 7, 4, 2$

$Z[4] = 8, 5$

$Z[5]_m = 9, 3, 1$

$Z[6]_a = 10, 7, 4, 2$

$Z[7] = 8, 5$

$Z[8]_m = 9, 3, 1$

$Z[9]_a = 10, 7, 4, 2$

$Z[10]_m = 9, 3, 1$

$Z[11]_u = \emptyset$

$Z = 9, 3, 1, 10, 7, 4, 2, 8, 5, 9, 3, 1, 10, 7, 4, 2, 8, 5, 9, 3, 1, 10, 7, 4, 2, 9, 3, 1$

Convert to another problem

Lemma: $\text{lcs}(S,T) = \text{lis}(Z)$

$\text{lis}(Z)$: length of the longest increasing subsequence of Z

$Z_{i_1} < Z_{i_2} < \dots < Z_{i_k}$ where $1 \leq i_1 < i_2 < \dots < i_k \leq n$

$\text{lis}(Z)$ contains j from $Z[i] \iff \text{lcs}(S,T)$ contains $S[i] = T[j]$

Example:

1 2 3 4 5 6 7 8 9 0 1

S: e m a _ m a _ m a m u e m a _ m a _ _ _ m a m u

T: m a m a _ s a _ m a - m a - m a _ s a _ m a - -

; 9, 3, 1; 10, 7, 4, 2; 8, 5; 9, 3, 1; 10, 7, 4, 2; 8, 5; 9, 3, 1; 10, 7, 4, 2; 9, 3, 1;

Problems

Let $r = |Z|$

- How to compute Z efficiently? (as a function of m, n, r, σ)
 - Small alphabet (e.g. $\Sigma = \{1, \dots, n + m\}$)
 - Large alphabet
 - Symbols = lines in files, as in diff
- How to find $\text{lis}(Z)$ efficiently?

A simple dynamic programming algorithm for lis(Z)

$A[i] = \text{l.i.s. of } Z_1 \dots Z_i \text{ that ends with } Z_i$

Example:

$Z = 9, 3, 1, 10, 7, 4, 2, 8, 5, 9, 3, 1 \dots$

$A = 1, 1, 1, 2, 2, 2, 2, 3, 3, 4, 3, 1, \dots$

Another dynamic programming algorithm for lis(Z)

Alg 2: $A[i, j] = \min. x$ such that x can be the last element in an increasing subsequence of $Z_1 \dots Z_i$ of length j

Example:

$i = 9, Z = 9, 3, 1, 10, 7, 4, 2, 8, 5, \dots$

$$A[9, 0] = -\infty$$

$$A[9, 1] = 1 \quad \text{i.s. (1)}$$

$$A[9, 2] = 2 \quad \text{i.s. (1, 2)}$$

$$A[9, 3] = 5 \quad \text{i.s. (1, 2, 5)}$$

$$A[9, i] = \infty \text{ for } i \geq 4$$

```

1  A[0,0] = -infinity; A[1..r] = infinity;
2  for(int i=1; i<=r; i++) {
3      A[i,0] = -infinity; A[0,i] = infinity;
4  }
5  for(int i=1; i<=r; i++) {
6      for(int j=1; j<=r; j++) {
7          if (A[i-1,j-1] < Z[i])
8              A[i,j] = min(Z[i], A[i-1,j]);
9          else
10             A[i,j] = A[i-1,j]
11     }
12 }
13 return highest k s.t. A[r,k]<infinity;

```

```
1 A[0] = -infinity; A[1..r] = infinity;
2 for(int i=1; i<=r; i++) {
3     find smallest k s.t. A[k]>=Z[i] (bin. search);
4     A[k] = Z[i];
5 }
6 return highest k s.t. A[k]<infinity;
```

```

1  A[0] = -infinity; A[1..r] = infinity;
2  B[0] = 0;
3  for(int i=1; i<=r; i++) {
4      find smallest k s.t. A[k]>=Z[i] (bin. search);
5      A[k] = Z[i];
6      B[k] = new node;
7      B[k]—>index = i; B[k]—>next = B[k-1];
8  }
9  find highest k s.t. A[k]<infinity;
10 v = B[k];
11 while(v) {
12     print v—>index; v = v—>next;
13 }

```