# Algorithmns and Datastructures Levenshtein distance, Dynamic programming

Albert-Ludwigs-Universität Freiburg

Prof. Dr. Rolf Backofen

Bioinformatics Group / Department of Computer Science Algorithmns and Datastructures, February 2017

# Structure



Introduction

Edit distance

- Measurement for similarity of two words / strings
- Algorithm for efficient calculation
- General principle: dynamic programming







#### Ergebnisse für eyjafjallajökull

Stattdessen suchen nach: ejafjatlajökuk

#### Eyjafjallajökull - Wikipedia

de.wikipedia.org/wiki/Eyjafjallajökull -

Der Name Eyjafjallajökull (isländisch für "Inselberge-Gletscher") rührt von den so genannten Landeyjar (dt. Landinseln) her. Das sind felsige Erhebungen, ... Name - Der Gletscher - Der Vulkan unter dem Gletscher - Eruptionsgeschichte

## Eyjafjallajökull - Der unaussprechliche Vulkanfilm Film 2014 ... www.kino.de > Filme 🔻

31.07.2014 - **Eyjafjallajökull** - Der unaussprechliche Vulkanfilm, Irrwitzige Komödie um ein verfeindetes Ex-Ehepaar, das wegen der Asche des isländischen ...

#### Bilder zu eyjafjallajökull

Unangemessene Bilder melden



Weitere Bilder zu eyjafjallajökull



### Eyjafjallajökull

Gletscher in Island

Der Eyjafjallajökull, zu deutsch Eyjafjöll-Gletscher, ist der sechstgrößte Gletscher Islands. Er liegt an der äußersten Südküste, westlich des Gletschers Mýrdalsjökull in der Gemeinde Rangárþing eystra, die größte Höhe beträgt 1651 m. Wikipedia

Letzte Eruption: April 2010

Höhe: 1.666 m Fläche: 100 km²

Prominenz: 1.051 m Erstbesteiger: Sveinn Pálsson

# A lot of applications where similar string are searched:

Duplicates in databases:

```
Hein Blöd 27568 Bremerhaven
Hein Bloed 27568 Bremerhafen
Hein Doof 27478 Cuxhaven
```

Product search:

memory stik

Websearch:

```
eyjaföllajaküll
uniwersität verien 2017
```

Bioinformatics: Similarity of DNA-sequences

# Search of similar proteins:

- BLAST (Basic Local Alignment Search Tool)
- Alignment ê Edit distance
- Changed life-science completely

# Google-Scholar entry:

[нтмь] Gapped **BLAST** and PSI-**BLAST**: a new generation of protein database search programs

<u>SF Altschul, TL Madden, AA Schäffer...</u> - Nucleic acids ..., 1997 - Oxford Univ Press Abstract The **BLAST** programs are widely used tools for searching protein and DNA databases for sequence similarities. For protein comparisons, a variety of definitional, algorithmic and statistical refinements described here permits the execution time of the ... Zitiert von: 58805 Ahnliche Artikel Alle 135 Versionen Zitieren Speichern

# **Definition of edit distance**: (Levenshtein-distance)

- Let x, y be two strings
- Edit distance ED(x,y) of x and y:
  The minimal number of operations to transform x into y
  - Insert a character
  - Replace a character with another
  - Delete a character

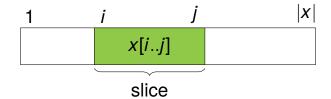
Example



```
12345
                           12345
                           BLOED
DOOF
        replace(1, B)
                                    replace(5, F)
                           BLOEF
BOOF
        replace(2, L)
                                    delete(4)
BLOF
                           BLOF
        insert(4, E)
                                    replace(2, 0)
BLOEF
                           BOOF
        replace(5, D)
                                    replace(1, D)
BLOED
                           DOOF
              ED=4
                                         ED=4
```

### **Notation:**

- lacksquare  $\epsilon$  is the empty string
- |x| is the length of the string x (number of characters)
- x[i..j] is the slice of x from i to j where  $1 \le i \le j \le |x|$



### Trivial facts:

- $\blacksquare$  ED(x, y) = ED(y, x)
- $\blacksquare$  ED( $x, \varepsilon$ ) = |x|

■ ED
$$(x,y) \ge abs(|x|-|y|)$$
 abs $(x) = \begin{cases} x & \text{if } x \ge 0 \\ -x & \text{else} \end{cases}$ 

■ 
$$ED(x,y) \le ED(x[1..n-1],y[1..m-1]) + 1$$
  $n = |x|, m = |y|$ 

# Solutions based on examples:

- From VERIEN to FERIEN?
- From MEXIKO to AMERIKA?
- From AAEBEAABEAREEAEBA to RBEAAEEBAAAEBBAEAE?
- Searching biggest substrings can yield the solution but doesn't have to

# Recursive approach:

Dividing in two halves? Not a good idea:

$$ED(GRAU, RAUM) = 2$$
 but  $ED(GR, RA) + ED(AU, UM) = 4$ 

Finding "smaller" sub problems? Let's try it!

# **Terminology:**

- $\blacksquare$  Let x, y be two strings
- Let  $\sigma_1, ..., \sigma_k$  be a sequence of k operations where  $k = \mathrm{ED}(x, y)$  for  $x \to y$  (transform x into y)

  (We do not know this sequence but we assume it exists)



## Terminology:

■ We only consider monotonous sequences: The positition of  $\sigma_{i+1}$  is  $\geq$  the position of  $\sigma_i$  where we only allow the positions to be equal on a delete operation

```
12345
                          1234567
                          SAUDOOF
DOOF
        replace(1, B)
                                      delete(1)
                          AUDOOF
BOOF
        replace(2, L)
                                      delete(1)
BLOF
                          UDOOF
        insert(4, E)
                                      delete(1)
BLOEF
                          DOOF
        replace(5, D)
                                      insert(4, 0)
BI OFD
```

# Terminology:

- **Lemma:** For any x and y with k = ED(x,y) exists a monotonous sequence of k operations for  $x \to y$
- Intuition: The order of our sequence is not relevant (Therefore we can also sort them monotonously)

# Consider the last operation:

■ Solve blue part recursively

DOOF ↓↓↓↓ BLOE ↓insert BLOED

Figure: Case 1a

DOOF  $\downarrow \downarrow \downarrow \downarrow \downarrow \downarrow \downarrow$ BLOEDF  $\downarrow \text{delete}$ BLOED

Figure: Case 1b

DOOF ↓↓↓↓↓ BLOEF ↓replace

BLOED

DLOLD

Figure: Case 1c

# Consider the last operation:

■ Solve blue part recursively

WINTER  $\downarrow \downarrow \downarrow \downarrow \downarrow \downarrow \downarrow$  SOMMER  $\downarrow nothing$  SOMMER

Figure: Case 2

# Display of solution:

- Alignment
- Example:



- Instances of Bellman's principle of optimality:
  - Shortest paths
  - Edit distance



Figure: Richard Bellman (1920 - 1984)

- Optimal solutions consist of optimal partial solutions
  - Shortest paths: Each partial path has to be optimal
  - Edit distance: Each partial alignment has to be optimal

 Always solvable through dynamic programming (Caching of optimal partial solutions)

# Case analysis:

- We consider the last operation  $\sigma_k$ 
  - $\sigma_1, ..., \sigma_{k-1}$ :  $x \to z$  and  $\sigma_k$ :  $z \to y$  Example:

$$x = DOOF$$
,  $z = SAUBLOEF$ ,  $y = SAUBLOED$ 

- Let n = |x|, m = |y|, m' = |z|
- We note  $m' \in \{m-1, m, m+1\}$  why?

# Case analysis:

■ Case 1:  $\sigma_k$  does something at the outer end:

```
■ Case 1a: \sigma_k = \text{insert}(m'+1, y[m]) [then m' = m-1]
■ Case 1b: \sigma_k = \text{delete}(m') [then m' = m+1]
■ Case 1c: \sigma_k = \text{replace}(m', y[m]) [then m' = m]
```

■ Case 2:  $\sigma_k$  does nothing at the outer end:

```
■ Then z[m'] = y[m] and x[n'] = z[m'] and with that \sigma_1, ..., \sigma_{k-1} : x[1..n-1] \rightarrow y[1..m-1] and x[n] = y[m]
```



# Case analysis:

- Case 1a (insert):  $\sigma_1, \ldots, \sigma_{k-1}$ :  $X \rightarrow y[1..m-1]$
- Case 1b (delete):  $\sigma_1, ..., \sigma_{k-1}$ :  $x[1..n-1] \rightarrow y$
- Case 1c (replace):  $\sigma_1, \ldots, \sigma_{k-1}$ :  $x[1..n-1] \rightarrow y[1..m-1]$
- Case 2 (nothing):  $\sigma_1, \ldots, \sigma_k$ :  $x[1..n-1] \rightarrow y[1..m-1]$

### This results in the recursive formula:

- For |x| > 0 and |y| > 0 is ED(x, y) the minimum of
  - ED(x , y[1..m-1]) + 1 and
  - $\blacksquare$  ED(x[1..n-1],y ) + 1 and
  - ED(x[1..n-1],y[1..m-1])+1 if  $x[n] \neq y[m]$
  - $\blacksquare$  ED(x[1..n-1], y[1..m-1]) + 0 if x[n] = y[m]
- For |x| = 0 is ED(x, y) = |y|
- For |y| = 0 is ED(x, y) = |x|

```
def edit_distance(x, y):
    if len(x) == 0:
        return len(y)
    if len(y) == 0:
        return len(x)
    ed1 = edit distance(x, y[:-1]) + 1
    ed2 = edit distance(x[:-1], y) + 1
    ed3 = edit_distance(x[:-1], y[:-1])
    if x[-1] != v[-1]:
        ed3 += 1
    return min(ed1, ed2, ed3)
```

■ The algorithm results in the following recursive formular:

$$T(n,m) = T(n-1,m) + T(n,m-1) + T(n-1,m-1) + 1$$

$$\geq T(n-1,m-1) + T(n-1,m-1) + T(n-1,m-1)$$

$$= 3 \cdot T(n-1,m-1)$$

- This results in  $T(n,n) \ge 3^n$
- ⇒ The runtime is at least exponential

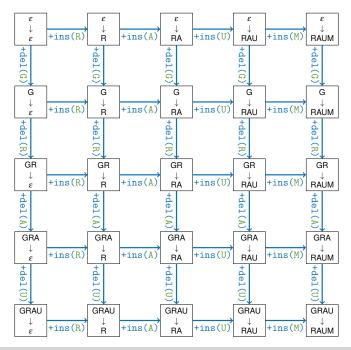
# Dynamic programming:

- We create a table with all possible combination of substrings and save calculated entries
- This results in a runtime and space consumption of  $O(n \cdot m)$

### Visualization on the next slide:

- Operations always refer to the last position (indices are omitted)
- We also display the replaced character on a replace operation to visualize operations without costs

$$\Rightarrow$$
 repl(A, A)

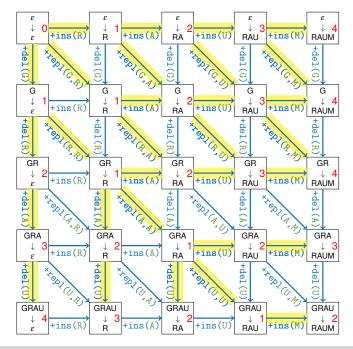


Fast algorithm



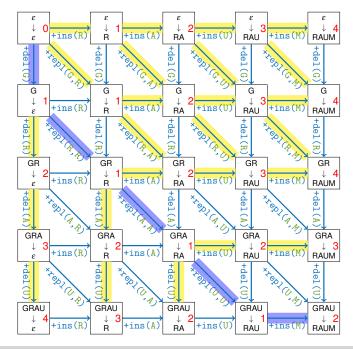
# Fast algorithm:

We can determine the edit distance for all combination of partial strings from the top left to bottom right.



# How to get the sequence of operations?

- We save at each recursion the most efficient previous entry (the highlighted arrows in our image)
- There can be more than one arrows to the three previous entries
- If we follow the highlighted path from (n,m) to (1,1) we get the optimum operations to transform x into y
  - If we can follow more than one path there exist more than one ideal sequence



# General principle:

- Recursive computation of ...
  - ... the same reoccuring partial problems
  - ... a limited number of partial problems
- Computation of the solutions for all partial problems
- In a order that unsolved partial problems consist of already solved partial problems
- The "path" to our solution normally gets computed while searching the best solution
- Dijkstra algorithm is basically dynamic programming!

# Additional applications:

- Edit distance: global alignment with  $O(n^2)$  space and time consumption
- But: Model for deletition unrealistic
  - In evolution larger pieces are more likely
  - delete operation: first gap expensive (e.g. 2), remainding are cheaper (e.g. 0.5)

Solution in  $O(n^3)$  time or  $O(n^2)$  affine

Additional applications (II)



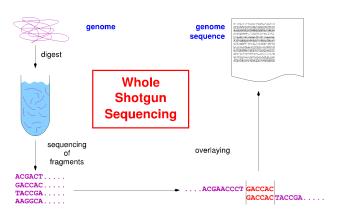
 $O(n^2)$  space consumption might be problematic:

# Hirschberg algorithm:

- Divide-and-conquer approach
- O(n) space and  $O(n^2)$  time consumption

### Additional applications (III)





- Sequencing:  $O(n^2)$  is too much
- Index: suffixtree, suffixarray, burrow-wheeler-transform

### ■ General

- [CRL01] Thomas H. Cormen, Ronald L. Rivest, and Charles E. Leiserson. Introduction to Algorithms. MIT Press, Cambridge, Mass, 2001.
- [MS08] Kurt Mehlhorn and Peter Sanders. Algorithms and data structures, 2008. https://people.mpi-inf.mpg.de/~mehlhorn/ftp/Mehlhorn-Sanders-Toolbox.pdf.

# Dynamic programming

```
[Wik] Dynamic programming
    https:
    //en.wikipedia.org/wiki/Dynamic_programming
```

### Edit distance

```
[Wik] Levenshtein distance
    https:
    //en.wikipedia.org/wiki/Levenshtein_distance
```