### Similarity Search in Large Databases Introduction to Similarity Search

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### What is Similarity Search?

• Similarity search deals with the question:

How similar are two objects?

- "Objects" may be
  - strings (Augsten ↔ Augusten)
  - tuples in a relational database

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- documents (e.g., HTML or XML)

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• "Similar" is application dependant

# Outline



- Intuition
- Applications
- Framework

Similarity Search Applications

Similarity Search

### Application I: Object Identification

- Problem:
  - Two data items represent the same real world object (e.g., the same
  - but they are represented differently in the database(s).
- How can this happen?
  - different coding conventions (e.g., Gilmstrasse, Hermann-von-Gilm-Str.)
  - spelling mistakes (e.g., Untervigil, Untervigli)
  - outdated values (e.g., Siegesplatz used to be Friedensplatz).
  - incomplete/incorrect values (e.g., missing or wrong apartment number in residential address).
- Focus in this course!

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Similarity Search Applications

#### Application I: Flavors of Object Identification

#### Duplicate Detection

- one table
- find all tuples in the table that represent the same thing in the real
- Example: Two companies merge and must build a single customer database.

#### Similarity Join

- two tables
- join all tuples with similar values in the join attributes
- Example: In order to detect tax fraud, data from different databases need to be linked.

#### Similarity Lookup

- one table, one tuple
- find the tuple in the table that matches the given tuple best
- Example: Do we already have customer X in the database?

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## Application III: Error Correction in Signal Processing

- Application: Transmit text signal over physical channel
- Problem: Transmission may introduce errors
- Goal: Restore original (sent) message
- Solution: Find correct text that is closest to received message.

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### Application II: Computational Biology

- DNA and protein sequences
  - modelled as text over alphabet (e.g.  $\{A, C, G, T\}$  in DNA)
- Application: Search for a pattern in the text
  - look for given feature in DNA
  - compare two DNAs
  - decode DNA
- Problem: Exact matches fail
  - experimental measures have errors
  - small changes that are not relevant
  - mutations
- Solution: Similarity search
  - Search for *similar* patterns
  - How similar are the patterns that you found?

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### Framework for Similarity Search

- 1. Preprocessing (e.g., lowercase Augsten  $\rightarrow$  augsten)
- 2. Search Space Reduction
  - Blocking
  - Sorted-Neighborhood
  - Filtering (Pruning)
- 3. Compute Distances
- 4. Find Matches

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### Search Space Reduction: Brute Force

- We consider the example of similarity join.
- Similarity Join: Find all pairs of similar tuples in tables A and B.
  - Search space:  $A \times B$  (all possible pairs of tuples)
  - Complexity: compute |A||B| distances  $\rightarrow$  expensive!  $(|A| = 30k, |B| = 40k, 1ms \text{ per distance} \Rightarrow \text{ join runs 2 weeks})$

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• Example: 16 distance computations!

Α		В		
Tim	m	Bil	m	
Bill	m	Jane	f	
Jane	f	Tim	m	
Mary	f	Marie	f	

• Goal: Reduce search space!

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### Search Space Reduction: Sorted Neighborhood

- Sorted Neighborhood
  - Sort A and B (e.g., by one of the attributes).
  - Move a window of fixed size over A and B.
    - move A-window if sort attribute of next tuple in A is smaller than in B
    - otherwise move B-window
  - Compare only tuples within the windows.
- Example: Sort by name, use window of size 2:

Α			В	
	Bill	m	Bil	m
	Jane	f	<b>J</b> ane	f
	Mary	f	<b>→</b> Marie	f
	Tim	m	Tim	m

• Improvement: 12 distance computations (instead of 16)!

### Search Space Reduction: Blocking

- Blocking
  - Partition A and B into blocks (e.g., group by chosen attribute).

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- Compare only tuples within blocks.
- Example: Block by gender (m/f):

Tim	m Bil	m
Bill	Tim	m

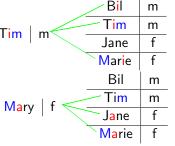
Mary	f Jane	f
Jane	f Marie	f

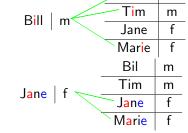
• Improvement: 8 distance computations (instead of 16)!

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### Search Space Reduction: Filtering

- Filtering (Pruning)
  - Remove (filter) tuples that cannot match, then compute the distances.
  - Idea: filter is faster than distance function.
- Example: Do not match names that have no character in common:





• Improvement: 11 distance computations (instead of 16)!

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#### Distance Computation

#### Definition (Distance Function)

Given two sets of objects, A and B, a distance function for A and B maps each pair  $(a, b) \in A \times B$  to a positive real number (including zero).

$$\delta: A \times B \to \mathbb{R}_0^+$$

- We will define distance functions for
  - sets
  - strings
  - ordered, labeled trees
  - unordered, labeled trees

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# Finding Matches: Threshold

$$\begin{array}{c|ccccc} & b_1 & b_2 & b_3 \\ \hline a_1 & 6 & 5 & 4 \\ a_2 & 2 & 2 & 1 \\ a_3 & 1 & 3 & 0 \\ \end{array}$$

- Once we know the distances which objects match?
- Threshold Approach:
  - fix threshold  $\tau$
  - algorithm:

foreach  $d_{ii} \in D$  do if  $d_{ii} < \tau$  then match  $(a_i, b_i)$ 

- produces *n*:*m*-matches
- Example with  $\tau = 3$ :  $\{(a_2, b_1), (a_2, b_2), (a_2, b_3), (a_3, b_1), (a_3, b_3)\}$

#### Distance Matrix

#### Definition (Distance Matrix)

Given a distance function  $\delta$  for two sets of objects,  $A = \{a_1, \dots, a_n\}$  and  $B = \{b_1, \ldots, b_m\}.$ 

The distance matrix D is an  $n \times m$ -matrix with

$$d_{ij} = \delta(a_i, b_i),$$

where  $d_{ii}$  is the element at the *i*-th row and the *j*-th column of D.

• Example distance matrix,  $A = \{a_1, a_2, a_3\}, B = \{b_1, b_2, b_3\}$ :

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Finding Matches: Global Greedy

Global Greedy Approach:

angularithm: 
$$M \leftarrow \emptyset$$
  $A \leftarrow \{a_1, a_2, \dots, a_n\}; B \leftarrow \{b_1, b_2, \dots, b_m\}$  create sorted list  $L$  with all  $d_{ij} \in D$  while  $A \neq \emptyset$  and  $B \neq \emptyset$  do  $d_{ij} \leftarrow$  deque smallest element from  $L$ 

if  $a_i \in A$  and  $b_i \in B$  then  $M \leftarrow M \cup (a_i, b_i)$ 

remove  $a_i$  from A and  $b_i$  from Breturn M

- produces 1:1-matches
- must deal with tie distances when sorting L! (e.g. sort randomly, sort by i and i)
- Example (sort ties by i, j):  $\{(a_3,b_3),(a_2,b_1),(a_1,b_2)\}$

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# Overview: Finding Matches

	$b_1$	$b_2$	$b_3$
$a_1$	6	5	4
$a_2$	2	2	1
<i>a</i> <sub>3</sub>	1	3	0

- Threshold Approach:
  - ullet all objects with distance below au match
  - produces *n*:*m*-matches
  - threshold approach for our example with  $\tau = 3$ :  $\{(a_2,b_1),(a_2,b_2),(a_2,b_3),(a_3,b_1),(a_3,b_3)\}$
- Global Greedy Approach:

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- pair with smallest distance is chosen first
- produces 1:1-matches
- global greedy approach for our example:  $\{(a_3,b_3),(a_2,b_1),(a_1,b_2)\}$

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Conclusion

- Framework for similarity queries:
  - 1. preprocessing
  - 2. search space reduction
    - blocking
    - sorted-neighborhood
    - filtering (pruning)
  - 3. compute distances: when are two objects similar?

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4. find matches: threshold, global greedy