Similarity Search in Large Databases Introduction to Similarity Search

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Similarity Search in Large Databases

Similarity Search Intuition

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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)
- "Similar" is application dependant

Similarity Search Outline Similarity Search Intuition Applications Framework

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Similarity Search Applications

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Application I: Entity Resolution (ER)

- Problem: (also known as "object identification")
 - Two data items represent the same real world object/entity (e.g., the same person),
 - 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 - outdated values (e.g., Siegesplatz used to be Friedensplatz).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 Entity Resolution

- Duplicate Detection (also: "dirty ER")
 - 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 (also: "clean-clean ER")
 - 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.

Application II: Computational Biology

- DNA and protein sequences
 - modelled as text over alphabet (e.g. $\{A, C, G, T\}$ in DNA)

Similarity Search Applications

- 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?

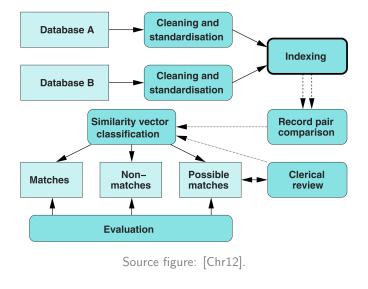
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Framework for Similarity Search / 1



Framework for Similarity Search / 2

1. Preprocessing: Cleaning and standardization, e.g.

• lowercase all values: Augsten \rightarrow augsten

• standardize values: $\{f, w, female, weiblich\} \rightarrow \{f\}$

• standardize encoding: K.Wolf Strasse → Karl-Wolf-Str.

2. Indexing for Search Space Reduction

blocking

sorted-neighborhood

• filter ing (pruning)

nearest neighbor search

3. Compute Distances

• compare record/tuple pairs

4. Find Matches: Classification

classify record/tuple pairs based on a distance or a vector of distances

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Search Space Reduction: Blocking

- Blocking
 - Generate blocks for A and B (seperately):
 - 1. compute one or more block keys for each tuple
 - 2. each distinct block key represents one block
 - 3. assign each tuple to all blocks of its block keys
 - Compare only tuples of block pairs with the same block key.
- Example: Attribute blocking: block key is value of a chosen attribute.

Block by "gender" attribute:

Tim	m Bil	m
Bill	Tim	m
Mary	fJane	f
Jane	f Marie	f

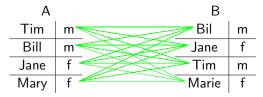
• Improvement: 8 distance computations (instead of 16)!

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.

Similarity Search Framework

- Search space: $A \times B$ (all possible pairs of tuples)
- Complexity: compute |A||B| distances \rightarrow expensive! $(|A| = 30k, |B| = 40k, 1ms \text{ per distance} \Rightarrow \text{ join will run for 2 weeks})$
- Example: 16 distance computations!



• Goal: Reduce search space!

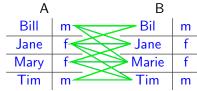
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Search Space Reduction: Sorted Neighborhood

- 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:



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

Search Space Reduction: Filtering

- Filtering (Pruning)
 - Filter: quick check if tuple pair is "promising" (i.e., distance is expected to be small enough).
 - Do not compute distance function if filter check fails.
 - Idea: filter is faster than distance function. Overhead for filtering is amortized by avoided distance computations.
- Example: Do not match names that have no character in common:

	Bil	m
Tim m	∕Tim	m
	Jane	f
	Marie	f
	Bil	m
Mary f	Tim	m
Ivialy I	Jane	f
	Marie	f

	Bil	m
Bill m	Tim	m
	Jane	f
	Marie	f
	Bil	m
Jane f	Tim	m
Jaile 1	Jane	f
•	Marie	f

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

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Distance Computation

Definition (Distance Function)

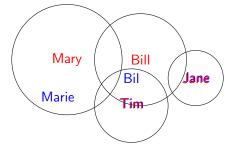
Given two sets of objects, A and B, a distance function for A and B maps each pair $(a, b) \in A \times B$ to a positive real number (including zero).

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

- We will define distance functions for
 - sets
 - strings
 - trees

Search Space Reduction: Nearest-Neighbor Search

- Represent tuples in some vector space or metric space.
- For each tuple t in A:
 - search for all tuples in B that are close to tuple t of A
 - distance to t is evaluated only for these tuples of B
- Example: Search for tuples of A in tuples of B.



• Improvement: 6 distance computations (instead of 16)!

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Distance Matrix

Definition (Distance Matrix)

Given a distance function δ 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

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

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

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

Finding Matches

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

- Once we know the distances which objects match?
- Distance matrix and search space reduction:
 - matrix may be only partially filled at the time of computing matches
 - missing distance values are treated as infinite

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Finding Matches: k-Nearest Neighbor Matching

- k-Nearest-Neighbor (kNN) Matching:
 - fix number of neighbors k
 - for a given object, form a match with its k nearest neighbors
- Algorithm: produces 1:k matching

foreach row *i* of distance matrix *D* **do** find column IDs c_1, c_2, \ldots, c_k of k smallest values in row i of Dform k matches $(a_i, b_{c_1}), (a_i, b_{c_2}), \dots, (a_i, b_{c_k})$

- Properties:
 - not symmetric: transposed matrix D^T may give a different result
 - ties affect the matching (only for k-th neighbor)
- Example k = 2: $\{(a_1, b_3), (a_1, b_2), (a_2, b_3), (a_2, b_1), (a_3, b_3), (a_3, b_1)\}$

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Finding Matches: Threshold Matching

- Threshold Approach:
 - fix threshold τ
 - ullet for a given object, all objects with a distance range of au are matched
- Algorithm: produces n:m matching

foreach
$$d_{ij} \in D$$
 do if $d_{ij} \le \tau$ then match (a_i, b_j)

• Example with $\tau = 2$: $\{(a_2, b_1), (a_2, b_2), (a_2, b_3), (a_3, b_1), (a_3, b_3)\}$

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Finding Matches: Global Greedy

- Global Greedy Approach
 - form object pair with smallest distance first
 - matched objects are removed

6 5 4

• Algorithm: produces 1:1 matching

$$M \leftarrow \emptyset$$

 $A \leftarrow \{a_1, a_2, \dots, a_n\}; \ B \leftarrow \{b_1, b_2, \dots, b_m\}$
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 $b_j \in B$ then
 $M \leftarrow M \cup (a_i, b_j)$
remove a_i from A and b_j from B

return M

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

Overview: Matching Techniques

- Threshold Matching:
 - ullet all objects with distance within au match
 - n:m-matching
 - symmetric, not affected by ties
- k-Nearest Neighbor Matching:
 - each object is matched to its k closest objects
 - 1:*k*-matching
 - not symmetric, affected by ties
- Global Greedy Approach:
 - pair with smallest distance is matched first and removed
 - 1:1-matching
 - symmetric, affected by ties

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Peter Christen.

A survey of indexing techniques for scalable record linkage and deduplication.

IEEE Trans. Knowl. Data Eng., 24(9):1537-1555, 2012.

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• Framework for similarity queries:

1. Preprocessing: Cleaning and standardization
2. Indexing for Search Space Reduction

• blocking
• sorted-neighborhood
• filtering (pruning)
• nearest neighbor search

3. Compute Distances

5. Compute Distances

4. Find Matches: Classification

threshold-based

• k-nearest-neighbor

• global greedy