Efficient Top-k Spatial Distance Joins

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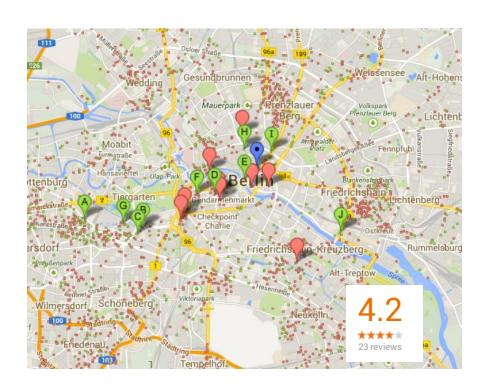
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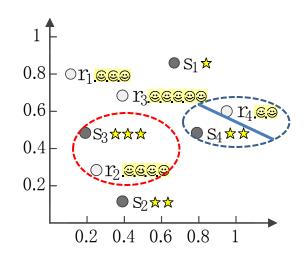
Motivation Example

- Observation
 - More spatial locations are tagged with score information (quality, rating, price, etc)
- Spatial object
 - Location: (x, y)
 - Score



Motivation Example

- Dataset R: restaurants
- Dataset S: hotels
- Recommend to tourists hotelrestaurant pairs, which are
 - (1) close to each other (spatial join)
 - (2) of high quality (top-k selection)



Example

$$k = 1, \epsilon = 0.2, SUM$$

Result

$$(r_2, s_3)$$

Definition

- Top-k Spatial Distance Joins (k-SDJ)
 - Two collections of spatial objects R and S (id, loc, score)
 - The k-SDJ query retrieves a k-subset of $R \times S$ such that for every result pair (r,s):
 - (1) r is spatially close to s (i.e., $dist(r,s) \le \epsilon$)
 - (2) it has top-k largest aggregation score $\gamma(r,s)$
 - $-\gamma$ is a monotone function

Contributions

- k-SDJ query
 - Location-based services
 - data integration
 - geoscience, etc.
- Algorithms
 - SFA, DFA, BA
- Experiments
 - Effectiveness and efficiency

Outline

- Related Work
 - Top-k Join Queries
 - Spatial Join Queries
- Algorithms
- Experiments
- Conclusions

Top-k Join Queries

- Rank aggregation on top of relational join results, i.e., assume simple join attributes and join predicate [Ilyas 2008]
- Hash-Based Rank-Join (HRJN) [Ilyas 2003]
 - Sort R and S descending to scores
 - HashTable indexes on R and S
 - Each accessed record is joined with the opposite hash table

Spatial Join Queries

- ϵ -Distance Join
 - Given two sets of spatial objects R and S, identify the object pairs (r,s) with $r \in R, s \in S$, such that $dist(r,s) \le \epsilon$
 - PlaneSweep Algorithm [Arge 1998]
 - For non-indexed inputs, sort and sweep
 - R-tree Join Algorithm [Brinkhoff 1993]
 - For inputs indexed by R-trees

Outline

- Related Work
- Algorithms
- Experiments
- Conclusions

k-SDJ Evaluation

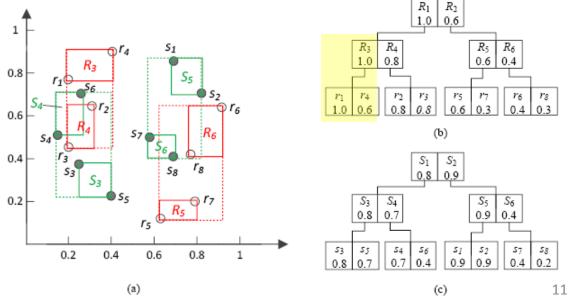
- Combination of top-k query and spatial join
 - Score: $\gamma(r,s)$
 - Distance: dist(r, s)
- Challenges
 - Effective pruning and termination techniques based on both dimensions
- Ideas
 - Index both spatial and score attributes
 - Score-first/Distance-first/Hybrid processing

Indexing: aR-tree

- R-tree augmented with maximum score information
- Property:
 - Given two aR-tree entries e_R and e_S . If $\gamma(e_R, e_S) = \beta$, $\forall r_i \in e_R$, $\forall s_i \in e_S$, $\gamma(r_i, s_i) \leq \beta$

| loc | score |
|--------------|--|
| (0.20, 0.78) | 1.0 |
| | 0.8 |
| (0.20, 0.45) | 0.8 |
| (0.40, 0.90) | 0.6 |
| (0.63, 0.12) | 0.6 |
| (0.91, 0.63) | 0.4 |
| (0.79, 0.20) | 0.3 |
| (0.76, 0.42) | 0.1 |
| | (0.20, 0.78) (0.30, 0.64) (0.20, 0.45) (0.40, 0.90) (0.63, 0.12) (0.91, 0.63) (0.79, 0.20) |

| id | loc | score |
|-----------------|-----------------|-------|
| s_1 | (0.69, 0.85) | 0.9 |
| s_2 | (0.81, 0.71) | 0.9 |
| s_3 | (0.24, 0.38) | 0.8 |
| s_4 | (0.15, 0.52) | 0.7 |
| s_5 | (0.40, 0.22) | 0.7 |
| s_6 | (0.25, 0.70) | 0.4 |
| s_7 | (0.58, 0.50) | 0.4 |
| s_8 | (0.68, 0.42) | 0.2 |
| \overline{Oh} | iect collection | on S |



- HRJN variant with aR-trees
 - Sorted according to scores
 - aR-tree indexes A_R and A_S on R/S
 - Each accessed record is
 - (1) Joined with the opposite aR-tree
 - (2) Inserted into corresponding aR-tree
 - Termination based on score bounds

Example

| id | loc | score |
|-------|--------------|-------|
| r_1 | (0.20, 0.78) | 1.0 |
| r_2 | (0.30, 0.64) | 0.8 |
| r_3 | (0.20, 0.45) | 0.8 |
| r_4 | (0.40, 0.90) | 0.6 |
| r_5 | (0.63, 0.12) | 0.6 |
| r_6 | (0.91, 0.63) | 0.4 |
| r_7 | (0.79, 0.20) | 0.3 |
| r_8 | (0.76, 0.42) | 0.1 |

Object collection R

| id | loc | score |
|-------|--------------|-------|
| s_1 | (0.69, 0.85) | 0.9 |
| s_2 | (0.81, 0.71) | 0.9 |
| s_3 | (0.24, 0.38) | 0.8 |
| s_4 | (0.15, 0.52) | 0.7 |
| s_5 | (0.40, 0.22) | 0.7 |
| s_6 | (0.25, 0.70) | 0.4 |
| s_7 | (0.58, 0.50) | 0.4 |
| s_8 | (0.68, 0.42) | 0.2 |
| OL | | - 0 |

Object collection S

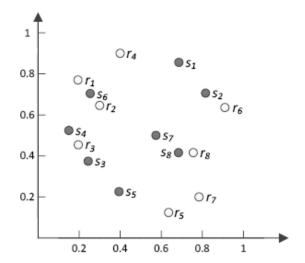
SFA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 1:

$$A_R = \{r_1\}$$
$$A_S = \{\}$$

$$A_{\mathcal{S}} = \{\}$$



Example

| id | loc | score |
|-----------------------|--------------|-------|
| r_1 | (0.20, 0.78) | 1.0 |
| r_2 | \ / | 0.8 |
| r_3 | (0.20, 0.45) | 0.8 |
| r_4 | (0.40, 0.90) | 0.6 |
| r_5 | (0.63, 0.12) | 0.6 |
| r_6 | (0.91, 0.63) | 0.4 |
| r_7 | (0.79, 0.20) | |
| r_8 | (0.76, 0.42) | 0.1 |
| Object collection R | | |

| id | loc | score |
|-------|--------------|-------|
| s_1 | (0.69, 0.85) | 0.9 |
| s_2 | (0.81, 0.71) | 0.9 |
| s_3 | (0.24, 0.38) | 0.8 |
| s_4 | (0.15, 0.52) | 0.7 |
| s_5 | (0.40, 0.22) | 0.7 |
| s_6 | (0.25, 0.70) | 0.4 |
| s_7 | (0.58, 0.50) | 0.4 |
| s_8 | (0.68, 0.42) | 0.2 |

Object collection S

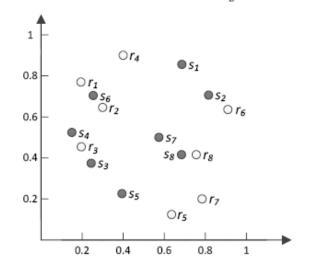
SFA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 2:

$$A_R = \{r_1\}$$
$$A_S = \{s_1\}$$

$$A_S = \{s_1\}$$

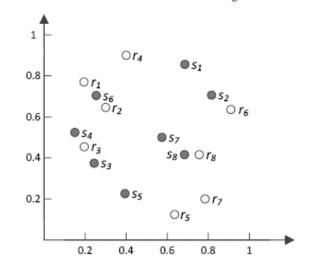


Example

| id | loc | score |
|-----------------------|--------------|-------|
| r_1 | (0.20, 0.78) | 1.0 |
| r_2 | (0.30, 0.64) | 0.8 |
| r_3 | (0.20, 0.45) | 0.8 |
| r_4 | (0.40, 0.90) | 0.6 |
| r_5 | (0.63, 0.12) | 0.6 |
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| Object collection R | | |

| | , | |
|-------|-----------------|--------|
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| s_7 | (0.58, 0.50) | 0.4 |
| s_8 | (0.68, 0.42) | 0.2 |
| Ob | ject collection | on S |

SFA Example: $k = 1, \epsilon = 0.1, \gamma = SUM$ Round 3: $A_R = \{r_1, r_2\}$ $A_S = \{s_1\}$



Example

| id | loc | score |
|-----------------------|--------------|-------|
| r_1 | (0.20, 0.78) | 1.0 |
| r_2 | (0.30, 0.64) | 0.8 |
| r_3 | (0.20, 0.45) | 0.8 |
| r_4 | (0.40, 0.90) | 0.6 |
| r_5 | (0.63, 0.12) | 0.6 |
| r_6 | (0.91, 0.63) | 0.4 |
| r_7 | (0.79, 0.20) | 0.3 |
| r_8 | (0.76, 0.42) | 0.1 |
| Object collection R | | |

| id | loc | score |
|-------|--------------|-------|
| s_1 | (0.69, 0.85) | 0.9 |
| s_2 | (0.81, 0.71) | 0.9 |
| s_3 | (0.24, 0.38) | 0.8 |
| s_4 | (0.15, 0.52) | 0.7 |
| s_5 | (0.40, 0.22) | 0.7 |
| s_6 | (0.25, 0.70) | 0.4 |
| s_7 | (0.58, 0.50) | 0.4 |
| s_8 | (0.68, 0.42) | 0.2 |

Object collection S

SFA Example:

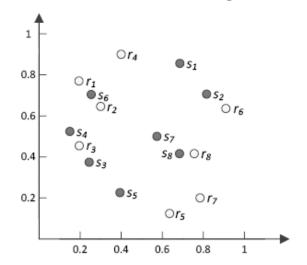
$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 4:

$$A_R = \{r_1, r_2\}$$

 $A_S = \{s_1, s_2\}$

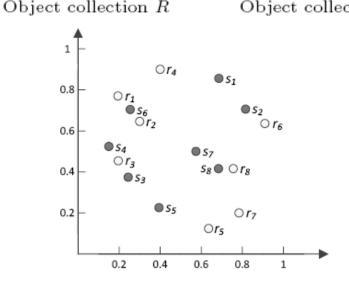
$$A_S = \{s_1, s_2\}$$



Example

| id | loc | score |
|-------|--------------|-------|
| r_1 | (0.20, 0.78) | 1.0 |
| r_2 | (0.30, 0.64) | 0.8 |
| r_3 | (0.20, 0.45) | 0.8 |
| r_4 | (0.40, 0.90) | 0.6 |
| r_5 | (0.63, 0.12) | 0.6 |
| r_6 | (0.91, 0.63) | 0.4 |
| r_7 | (0.79, 0.20) | 0.3 |
| r_8 | (0.76, 0.42) | 0.1 |

| id | loc | score |
|-------|--------------|-------|
| s_1 | (0.69, 0.85) | 0.9 |
| s_2 | (0.81, 0.71) | 0.9 |
| s_3 | (0.24, 0.38) | 0.8 |
| s_4 | (0.15, 0.52) | 0.7 |
| s_5 | (0.40, 0.22) | 0.7 |
| s_6 | (0.25, 0.70) | 0.4 |
| s_7 | (0.58, 0.50) | 0.4 |
| s_8 | (0.68, 0.42) | 0.2 |
| 88 | 3 | |



SFA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 6:

$$A_R = \{r_1, r_2, s_3\}$$

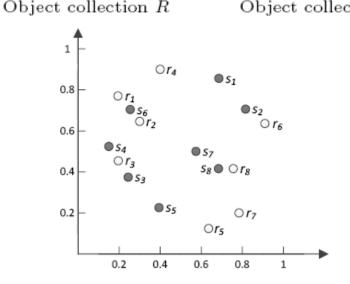
 $A_S = \{s_1, s_2, s_3\}$
 $C = \{(r_3, s_3) = 1.6\}$

Upper bound of remaining records: 1.8

Example

| id | loc | score |
|-------|--------------|-------|
| r_1 | (0.20, 0.78) | 1.0 |
| r_2 | (0.30, 0.64) | 0.8 |
| r_3 | (0.20, 0.45) | 0.8 |
| r_4 | (0.40, 0.90) | 0.6 |
| r_5 | (0.63, 0.12) | 0.6 |
| r_6 | (0.91, 0.63) | 0.4 |
| r_7 | (0.79, 0.20) | 0.3 |
| r_8 | (0.76, 0.42) | 0.1 |

| id | loc | score | |
|-----------------------|--------------|-------|--|
| s_1 | (0.69, 0.85) | 0.9 | |
| s_2 | (0.81, 0.71) | 0.9 | |
| s_3 | (0.24, 0.38) | 0.8 | |
| s_4 | (0.15, 0.52) | 0.7 | |
| 85 | (0.40, 0.22) | 0.7 | |
| s_6 | (0.25, 0.70) | 0.4 | |
| s_7 | (0.58, 0.50) | 0.4 | |
| s_8 | (0.68, 0.42) | 0.2 | |
| Object collection S | | | |



SFA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 10:

$$A_R = \{r_1, r_2, r_3, r_4\}$$

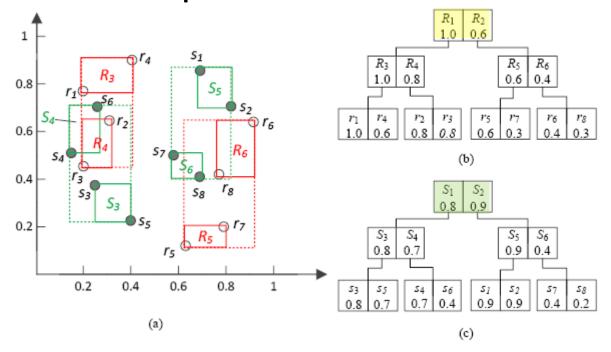
 $A_S = \{s_1, s_2, s_3, s_4, s_5, s_6\}$
 $C = \{(r_3, s_3) = 1.6\}$
Upper bound of remaining records: 1.5

return C

- Drawbacks:
 - Update operation on aR-trees
 - Range query operation for each record
 - Inefficient in spatial join computation

- General idea: Join -> Sort
- Challenge: Progressive join based on both spatial and aggregate score information
 - Bulk-load R and S to build aR-tree indexes A_R and A_S
 - Best-first order traversal
 - Use max-heap to prioritize aR-tree entry pairs based on $\gamma(e_R,e_S)$
 - Rationale: Entries containing objects with larger scores are processed first and thus enabling early termination

Example



DFA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

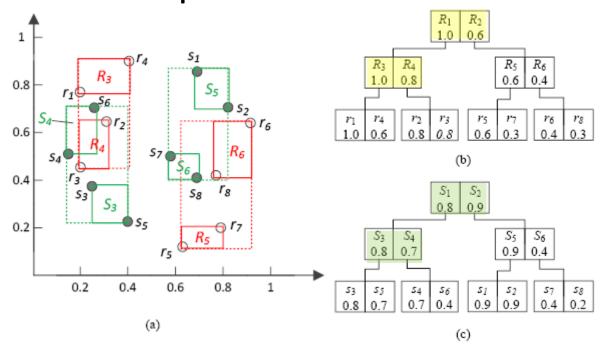
Round 1:

$$R_{root} \bowtie S_{root} = \{R_1S_1, R_2S_2\}$$

| H_e |
|-------------------|
| $(R_1, S_1), 1.8$ |
| $(R_2, S_2), 1.5$ |

 R_1S_2, R_2S_1 : pruned by distance

Example



DFA Example:

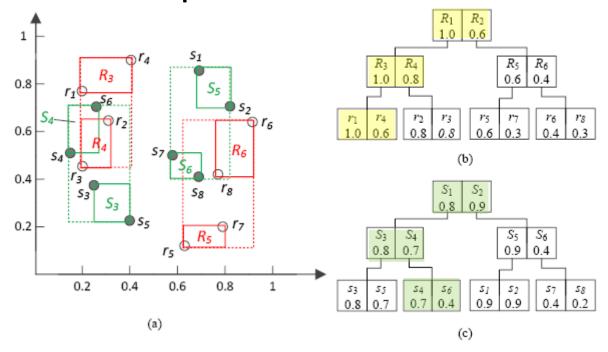
$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 2:

$$R_1 \bowtie S_1 = \{R_3S_4, R_4S_3, R_4S_4\}$$

| H_e | | |
|-------------------|--|--|
| $(R_3, S_4), 1.7$ | | |
| $(R_4, S_3), 1.6$ | | |
| $(R_4, S_4), 1.5$ | | |
| $(R_2, S_2), 1.5$ | | |

Example



DFA Example:

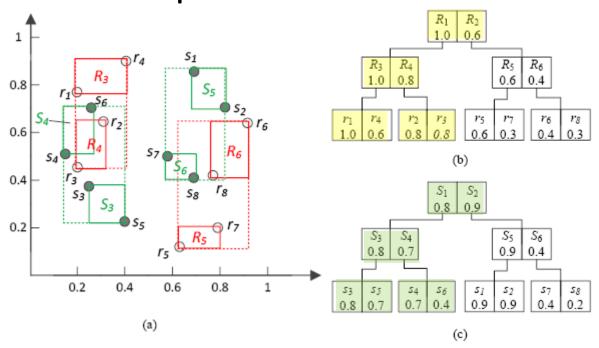
$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 3:

$$R_3 \bowtie S_4 = \{r_1 s_6\}$$

| H_e |
|-------------------|
| $(R_4, S_3), 1.6$ |
| $(R_4, S_4), 1.5$ |
| $(R_2, S_2), 1.5$ |
| $(r_1, s_6), 1.4$ |

Example



Given two aR-tree entries e_R and e_S . If $\gamma(e_R, e_S) = \beta$, $\forall r_i \in e_R$, $\forall s_j \in e_S$, $\gamma(r_i, s_j) \leq \beta$

DFA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 3:

$$R_4 \bowtie S_3 = \{r_3 s_3\}$$

| H_e |
|-------------------|
| $(r_3, s_3), 1.6$ |
| $(R_4, S_4), 1.5$ |
| $(R_2, S_2), 1.5$ |
| $(r_1, s_6), 1.4$ |

$$C = \{(r_3, s_3)\}$$
 return C

SFA vs. DFA

| | SFA | DFA | |
|---------|-------------------------------------|---------------------------------|--|
| Dataset | Score-ordered partial traversal | Index the whole datasets | |
| Index | Update the aR-trees for each record | Bulk-load to build the aR-trees | |
| Join | Probe the index for each record | Tree level join | |

- Challenge: exploit the benefits of SFA and DFA, while avoid their disadvantages
- Idea: Retrieve sorted records in block-wise fashion, bulk-load and join small blocks
 - Retrieve records in order of scores, thus able to terminate like SFA
 - Bulk-load aR-tree on each block for only once
 - Perform DFA on block level

Example

| | id | loc | score |
|----------|--------------|------------------------------|--------------|
| b_{R1} | $r_1 \\ r_2$ | (0.20, 0.78) (0.30, 0.64) | 1.0 0.8 |
| b_{R2} | r_3 r_4 | (0.20, 0.45) (0.40, 0.90) | 0.8 0.6 |
| b_{R3} | r_5 r_6 | (0.63, 0.12) (0.91, 0.63) | $0.6 \\ 0.4$ |
| b_{R4} | r_7 r_8 | (0.79, 0.20) (0.76, 0.42) | $0.3 \\ 0.1$ |
| | (a) | collection | R |

| | id | loc | score |
|----------|-------|------------------------------|-------|
| b_{S1} | s_1 | (0.69, 0.85) (0.81, 0.71) | 0.9 |
| 031 | s_2 | (0.81, 0.71) | 0.9 |
| b_{S2} | s_3 | (0.24, 0.38) | 0.8 |
| 052 | s_4 | (0.15, 0.52) | 0.7 |
| b_{S3} | s_5 | (0.40, 0.22) | 0.7 |
| 053 | s_6 | (0.25, 0.70) | 0.4 |
| har | s_7 | (0.58, 0.50) | 0.4 |
| b_{S4} | s_8 | (0.68, 0.42) | 0.2 |
| | (b) | collection | S |

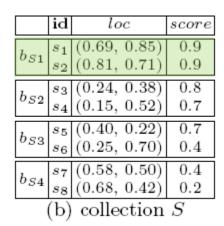
BA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 1: Build A_{R1} on block b_{R1} Join A_{R1} with \emptyset $C = \{\}$

Example

| | id | loc | score |
|----------|--------------|------------------------------|--------------|
| b_{R1} | $r_1 \\ r_2$ | (0.20, 0.78) (0.30, 0.64) | 1.0 0.8 |
| b_{R2} | r_3 r_4 | (0.20, 0.45) (0.40, 0.90) | 0.8 0.6 |
| b_{R3} | r_5 r_6 | (0.63, 0.12) (0.91, 0.63) | $0.6 \\ 0.4$ |
| b_{R4} | r_7 r_8 | (0.79, 0.20) (0.76, 0.42) | $0.3 \\ 0.1$ |
| | (a) | collection | R |



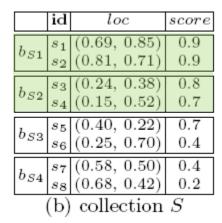
BA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 2: Build A_{S1} on block b_{S1} Join A_{S1} with A_{R1} $C = \{\}$

Example

| | id | loc | score |
|----------|--------------|------------------------------|--------------|
| b_{R1} | $r_1 \\ r_2$ | (0.20, 0.78) (0.30, 0.64) | 1.0 0.8 |
| b_{R2} | r_3 r_4 | (0.20, 0.45) (0.40, 0.90) | 0.8 0.6 |
| b_{R3} | r_5 r_6 | (0.63, 0.12) (0.91, 0.63) | $0.6 \\ 0.4$ |
| b_{R4} | r_7 r_8 | (0.79, 0.20) (0.76, 0.42) | $0.3 \\ 0.1$ |
| | (a) | collection | R |



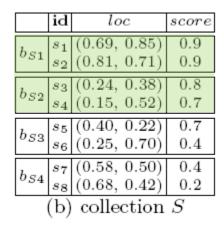
BA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 3: Build A_{S2} on block b_{S2} Join A_{S2} with A_{R1} $C = \{\}$

Example

| | id | loc | score |
|----------|-------------|------------------------------|--------------|
| b_{R1} | | (0.20, 0.78) (0.30, 0.64) | 1.0 0.8 |
| b_{R2} | r_3 r_4 | (0.20, 0.45) (0.40, 0.90) | 0.8 0.6 |
| b_{R3} | r_5 r_6 | (0.63, 0.12) (0.91, 0.63) | $0.6 \\ 0.4$ |
| b_{R4} | r_7 r_8 | (0.79, 0.20) (0.76, 0.42) | 0.3 0.1 |
| | (a) | collection | R |



BA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 4:

Build A_{R2} on block b_{R2} Join A_{R2} with A_{S1} and A_{S2} $C = \{(r_3, s_3)\}$

Example

| | id | loc | score | |
|------------------|--------------|------------------------------|--------------|--|
| b_{R1} | $r_1 \\ r_2$ | (0.20, 0.78) (0.30, 0.64) | 1.0 0.8 | |
| b_{R2} | r_3 r_4 | (0.20, 0.45) (0.40, 0.90) | 0.8 0.6 | |
| b_{R3} | r_5 r_6 | (0.63, 0.12) (0.91, 0.63) | $0.6 \\ 0.4$ | |
| b_{R4} | r_7 r_8 | (0.79, 0.20) (0.76, 0.42) | $0.3 \\ 0.1$ | |
| (a) collection R | | | | |

| | id | loc | score | |
|------------------|-------|------------------------------|-------|--|
| b_{S1} | s_1 | (0.69, 0.85) (0.81, 0.71) | 0.9 | |
| | s_2 | (0.81, 0.71) | 0.9 | |
| b_{S2} | s_3 | (0.24, 0.38) (0.15, 0.52) | 0.8 | |
| | s_4 | (0.15, 0.52) | 0.7 | |
| b_{S3} | s_5 | (0.40, 0.22) | 0.7 | |
| | s_6 | (0.40, 0.22) (0.25, 0.70) | 0.4 | |
| b_{S4} | s_7 | (0.58, 0.50) (0.68, 0.42) | 0.4 | |
| | s_8 | (0.68, 0.42) | 0.2 | |
| (b) collection S | | | | |

BA Example:

$$k = 1, \epsilon = 0.1, \gamma = SUM$$

Round 5:

Build A_{S3} on block b_{S3} Join A_{S3} with A_{R1} $C = \{(r_3, s_3) = 1.6\}$

Note:

 A_{S3} won't join with A_{R2} $\gamma(b_{R2}^u, b_{S3}^u) = 1.5 < 1.6$

$$l_R = 0.6, l_S = 0.4$$

 $T = 1.5 < \gamma(r_3, s_3)$
return C

- Access on block level
 - Avoid indexing whole datasets
- Bulk-load and join blocks
 - Avoid repeated update and range query
- Score bound for each block
 - Block-based pruning

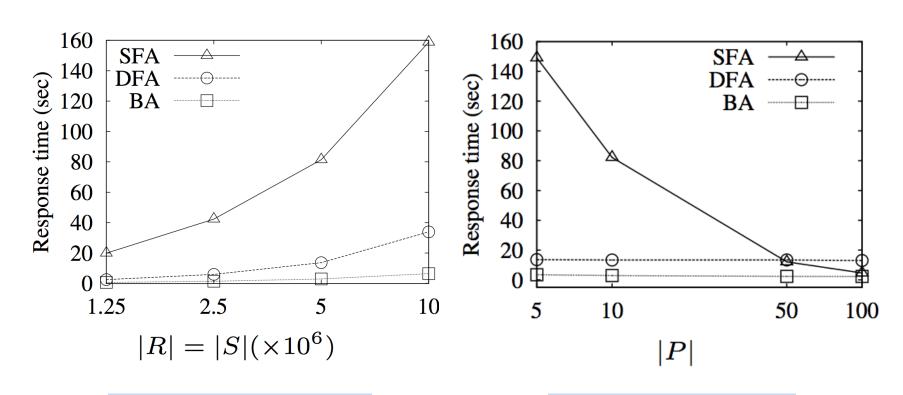
Outline

- Related Work
- Algorithms
- Experiments
- Conclusions

Experimental Evaluation

- Datasets
 - FLICKR, 1.68M locations in London
 - ISLES, 20M POIs from OpenStreetMap
 - Tag scores according to |P| seed points (simulating POIs) [Yiu 2011]
 - $o.score = 1 \min_{i=1}^{|P|} dist(o, P_i)$
- Assumptions
 - $-\gamma = SUM$
 - Data in memory

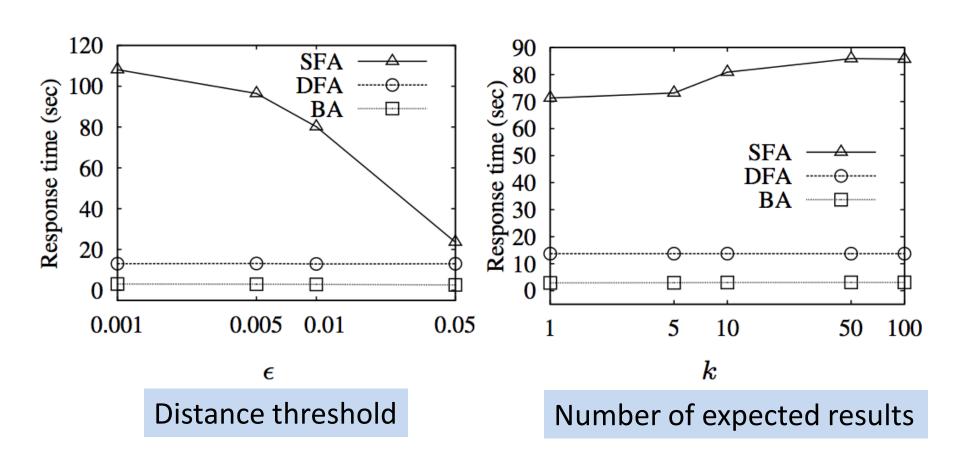
Data



Dataset cardinality

Number of seeds

Parameters



Outline

- Related Work
- Algorithms
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To sum up

Conclusions

- We proposed and studied the top-k spatial distance join query
- We proposed three k-SDJ solutions, i.e., SFA, DFA and BA
- According to our experiments, BA performs best in practice

Future work

- BA as general framework for complex top-k join queries (e.g., top-k string join)
- Theoretical analysis on optimal block size selection
- Combined distance/score ranking

Thank you

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