Efficient Parallel Partition based Algorithms for Similarity Search and Join with Edit Distance Constraints

Yu Jiang, Dong Deng, Jiannan Wang, Guoliang Li, and Jianhua Feng

Tsinghua University

Similarity Search&Join Competition on EDBT/ICDT 2013

Outline

Motivation

- Problem Definition
- Application
- Our Approach
 - Pass Join Algorithm
 - Additional Filters
 - Parallel

3 Experiment

- Evaluating Pruning Techniques
- Evaluating Parallelism
- Evaluating Scalability

Problem Definition Application

Problem Definition STRING SIMILARITY JOINS

Given a set of strings S, the task is to find all pairs of τ -similar strings from S. A program must output all matches with both string identifiers and distance τ .(Track II)

Problem Definition Application

An Example

Table: A string dataset

ID	Strings	Length
<i>S</i> 1	vankatesh	9
<i>s</i> ₂	avataresha	10
<i>s</i> ₃	kaushic chaduri	15
<i>S</i> 4	kaushik chakrab	15
S 5	kaushuk chadhui	15
<i>s</i> ₆	caushik chakrabar	17

Consider the string dataset in Table 1. Suppose $\tau = 3$. $\langle s_4, s_6 \rangle$ is a similar pair as $ED(s_4, s_6) \leq \tau$

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □

Problem Definition Application

Application

- Data cleaning
- Information Extraction
- Comparison of biological sequences

• ...

Pass Join Algorithm Additional Filters Parallel

Basic Idea

Lemma

Given a string r with τ + 1 segments and a string s, if s is similar to r within threshold τ , s must contain a segment of r.

Example

 $\tau = 1$, r = "EDBT" has two segments "ED" and "BT". s = "ICDT" cannot similar to r as s contains none of the two segements.

Pass Join Algorithm Additional Filters Parallel

Even Partition Scheme

Definition

In even partition scheme, each segment has almost the same length. $(\lfloor \frac{|s|}{\tau+1} \rfloor$ or $\lceil \frac{|s|}{\tau+1} \rceil)$

Example

 $\tau = 3$, we partition $s_1 =$ "vankatesh" into four segments "va", "nk", "at", "esh".

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □

Pass Join Algorithm Additional Filters Parallel

Substring Selection Basic Methods

- Enumeration: Enumerate all substrings for each of the segment.
- Length-based:

For each segment, only select substrings with same length.

• Shift-based:

For segment with start position p_i , select substrings with start position in $[p_i - \tau, p_i + \tau]$

Pass Join Algorithm Additional Filters Parallel

Substring Selection Position-aware Substring Selection

Observation

$$r_{l} \quad r_{r}$$

$$r = "vankatesh"$$

$$s = "avataresha"$$

$$s_{l} \quad s_{r}$$

$$||s_{l}| - |r_{l}|| + ||s_{r}| - |r_{r}|| = 2 + 3 > 3$$

Theorem (Position-aware Substring Selection)

For segment with start position p_i , select substrings with start position in $[p_i - \lfloor \frac{\tau - \Delta}{2} \rfloor, p_i + \lfloor \frac{\tau + \Delta}{2} \rfloor]$ where $\Delta = |s| - |r|$.

Pass Join Algorithm Additional Filters Parallel

Substring Selection Position-aware Substring Selection

Observation

$$r_{l} \quad r_{r}$$

$$r = "vankatesh"$$

$$s = "avataresha"$$

$$s_{l} \quad s_{r}$$

$$||s_{l}| - |r_{l}|| + ||s_{r}| - |r_{r}|| = 2 + 3 > 3$$

Theorem (Position-aware Substring Selection)

For segment with start position p_i , select substrings with start position in $[p_i - \lfloor \frac{\tau - \triangle}{2} \rfloor, p_i + \lfloor \frac{\tau + \triangle}{2} \rfloor]$ where $\triangle = |s| - |r|$.

Pass Join Algorithm Additional Filters Parallel

Substring Selection Position-aware Substring Selection

Example

$$r = \text{``vankatesh''} \quad s = \text{``avataresha''}$$

$$p_1=1, \text{ va} \longleftarrow [1,3]: \text{ av va at}$$

$$p_2=3, \text{ nk} \longleftarrow [2,5]: \text{ va at ta ar}$$

$$p_3=5, \text{ at} \longleftarrow [4,7]: \text{ ta ar re es}$$

$$p_4=7, \text{ esh} \longleftarrow [6,8]: \text{ res esh sha}$$

$$\tau = 3, \Delta = 1, [p_i - \lfloor \frac{\tau - \Delta}{2} \rfloor, p_i + \lfloor \frac{\tau + \Delta}{2} \rfloor] = [p_i - 1, p_i + 2]$$

Pass Join Algorithm Additional Filters Parallel

Substring Selection Multi-match-aware Substring Selection

Observation



r, has3 segments to detect, 2 errors allowed

There must be another matching between r_r and s_r .

Theorem (Multi-match-aware Substring Selection)

For the *i*-th segment with start position p_i , select substrings within $[p_i-i, p_i+i] \cap [p_i+\triangle - (\tau+1-i), p_i+\triangle + (\tau+1-i)].$

Pass Join Algorithm Additional Filters Parallel

Substring Selection Multi-match-aware Substring Selection

Observation



r, has3 segments to detect, 2 errors allowed

There must be another matching between r_r and s_r .

Theorem (Multi-match-aware Substring Selection)

For the *i*-th segment with start position p_i , select substrings within $[p_i-i, p_i+i] \cap [p_i+\triangle - (\tau+1-i), p_i+\triangle + (\tau+1-i)].$

Pass Join Algorithm Additional Filters Parallel

Substring Selection Multi-match-aware Substring Selection



Pass Join Algorithm Additional Filters Parallel

Substring Selection Theoretical Results

- The number of selected substrings by the multi-match-aware method is minimum.
- 2 For strings longer than $2 * (\tau + 1)$, our selection method is the only way to select minimum number of substrings.

Pass Join Algorithm Additional Filters Parallel

Substring Selection Experimental Results



Figure: Numbers of selected substrings

・ロト (周) (E) (E) (E) (E)

Pass Join Algorithm Additional Filters Parallel

Substring Selection Experimental Results



Figure: Elapsed time for generating substrings

・ロト (周) (E) (E) (E) (E)

Pass Join Algorithm Additional Filters Parallel

Verification Length-aware Verification



- Inspired by the position-aware substring selection.
- Save at least half computation than traditional dynamic method.
- Save even more using improved early termination.

三日 のへで

э

Pass Join Algorithm Additional Filters Parallel

Verification Length-aware Verification



- Inspired by the position-aware substring selection.
- Save at least half computation than traditional dynamic method.
- Save even more using improved early termination.

э

ELE DQC

Pass Join Algorithm Additional Filters Parallel

Verification Length-aware Verification



- Inspired by the position-aware substring selection.
- Save at least half computation than traditional dynamic method.
- Save even more using improved early termination.

Pass Join Algorithm Additional Filters Parallel

Verification Extension-based Verification



- Inspired by the multi-match-aware substring selection.
- Using tighter thresholds to verify the candidate pairs.
- Verify if $ED(r_r, s_r) \leq \tau + 1 i$ and $ED(r_l, s_l) \leq i 1$.

Pass Join Algorithm Additional Filters Parallel

Verification Extension-based Verification



- Inspired by the multi-match-aware substring selection.
- Using tighter thresholds to verify the candidate pairs.
- Verify if $ED(r_r, s_r) \le \tau + 1 i$ and $ED(r_l, s_l) \le i 1$.

Pass Join Algorithm Additional Filters Parallel

Verification Extension-based Verification



- Inspired by the multi-match-aware substring selection.
- Using tighter thresholds to verify the candidate pairs.
- Verify if $ED(r_r, s_r) \le \tau + 1 i$ and $ED(r_l, s_l) \le i 1$.

ELE DQC

Pass Join Algorithm Additional Filters Parallel

Verification Experimental Results



Figure: Elapsed time for verification

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □

Pass Join Algorithm Additional Filters Parallel

Additional Filters Effective Indexing Strategy

• Partition longer strings into segments.

- Select substrings from shorter strings.
- Longer segments decrease the possibility of matching.
- Thus decrease the number of candidates.

Pass Join Algorithm Additional Filters Parallel

- Partition longer strings into segments.
- Select substrings from shorter strings.
- Longer segments decrease the possibility of matching.
- Thus decrease the number of candidates.

Pass Join Algorithm Additional Filters Parallel

Additional Filters Effective Indexing Strategy

- Partition longer strings into segments.
- Select substrings from shorter strings.
- Longer segments decrease the possibility of matching.
- Thus decrease the number of candidates.

Pass Join Algorithm Additional Filters Parallel

Additional Filters Effective Indexing Strategy

- Partition longer strings into segments.
- Select substrings from shorter strings.
- Longer segments decrease the possibility of matching.
- Thus decrease the number of candidates.

Pass Join Algorithm Additional Filters Parallel

Additional Filters Content Filter

Observation

- Let H_r denote the character frequency vector of r.
- r = abyyyy, s = axyyyxy. $\mathcal{H}_r = \{\{a, 1\}, \{b, 1\}, \{y, 4\}\}, \mathcal{H}_s = \{\{a, 1\}, \{x, 3\}, \{y, 4\}\}$
- Let $\mathcal{H}_{\Delta} = |\mathcal{H}_r \mathcal{H}_s|$.
- $\mathcal{H}_{\bigtriangleup} = |\mathcal{H}_r \mathcal{H}_s| = ||\mathbf{1}| + |-\mathbf{3}|| = 4.$
- A deletion or insertion changes $\mathcal{H}_{\bigtriangleup}$ by 1 at most.
- An substitution changes $\mathcal{H}_{\bigtriangleup}$ by 2 at most.

Pass Join Algorithm Additional Filters Parallel

Additional Filters Content Filter

Observation

- Let \mathcal{H}_r denote the character frequency vector of r.
- r = "abyyyy", s = "axxyyyxy".
 H_r = {{a,1}, {b,1}, {y,4}}, H_s = {{a,1}, {x,3}, {y,4}}
- Let $\mathcal{H}_{\triangle} = |\mathcal{H}_r \mathcal{H}_s|$.
- $\mathcal{H}_{\bigtriangleup} = |\mathcal{H}_r \mathcal{H}_s| = ||\mathbf{1}| + |-\mathbf{3}|| = 4.$
- A deletion or insertion changes $\mathcal{H}_{\bigtriangleup}$ by 1 at most.
- An substitution changes $\mathcal{H}_{\bigtriangleup}$ by 2 at most.

Pass Join Algorithm Additional Filters Parallel

Additional Filters Content Filter

Observation

- Let \mathcal{H}_r denote the character frequency vector of r.
- r = "abyyyy", s = "axxyyyxy".
 H_r = {{a,1}, {b,1}, {y,4}}, H_s = {{a,1}, {x,3}, {y,4}}
- Let $\mathcal{H}_{\Delta} = |\mathcal{H}_r \mathcal{H}_s|$.
- $\mathcal{H}_{\bigtriangleup} = |\mathcal{H}_r \mathcal{H}_s| = ||\mathbf{1}| + |-\mathbf{3}|| = 4.$
- A deletion or insertion changes $\mathcal{H}_{\bigtriangleup}$ by 1 at most.
- An substitution changes \mathcal{H}_{\triangle} by 2 at most.

Pass Join Algorithm Additional Filters Parallel

Additional Filters Content Filter

Observation

- Let \mathcal{H}_r denote the character frequency vector of r.
- r = "abyyyy", s = "axxyyyxy".
 H_r = {{a,1}, {b,1}, {y,4}}, H_s = {{a,1}, {x,3}, {y,4}}
- Let $\mathcal{H}_{\Delta} = |\mathcal{H}_r \mathcal{H}_s|$.
- $\mathcal{H}_{\bigtriangleup} = |\mathcal{H}_r \mathcal{H}_s| = ||\mathbf{1}| + |-\mathbf{3}|| = 4.$
- A deletion or insertion changes $\mathcal{H}_{\bigtriangleup}$ by 1 at most.
- An substitution changes $\mathcal{H}_{\bigtriangleup}$ by 2 at most.

Pass Join Algorithm Additional Filters Parallel

Additional Filters

Observation

- At most τ edit operations, $\mathcal{H}_{\triangle} \leq 2\tau$.
- At most $\tau ||\mathbf{r}| |\mathbf{s}||$ substitutions, $\mathcal{H}_{\triangle} \leq 2\tau ||\mathbf{r}| |\mathbf{s}||$.
- Group symbols to improve the content-filter running time.
- Integrate the content filter with the extension-based verification.

Pass Join Algorithm Additional Filters Parallel

Additional Filters

Observation

- At most τ edit operations, $\mathcal{H}_{\triangle} \leq 2\tau$.
- At most $\tau ||\mathbf{r}| |\mathbf{s}||$ substitutions, $\mathcal{H}_{\triangle} \leq 2\tau ||\mathbf{r}| |\mathbf{s}||$.
- Group symbols to improve the content-filter running time.
- Integrate the content filter with the extension-based verification.

Pass Join Algorithm Additional Filters Parallel

Additional Filters

Observation

- At most τ edit operations, $\mathcal{H}_{\triangle} \leq 2\tau$.
- At most $\tau ||\mathbf{r}| |\mathbf{s}||$ substitutions, $\mathcal{H}_{\triangle} \leq 2\tau ||\mathbf{r}| |\mathbf{s}||$.
- Group symbols to improve the content-filter running time.
- Integrate the content filter with the extension-based verification.

Pass Join Algorithm Additional Filters Parallel

Additional Filters Content Filter

Observation

- At most τ edit operations, $\mathcal{H}_{\triangle} \leq 2\tau$.
- At most $\tau ||\mathbf{r}| |\mathbf{s}||$ substitutions, $\mathcal{H}_{\triangle} \leq 2\tau ||\mathbf{r}| |\mathbf{s}||$.
- Group symbols to improve the content-filter running time.
- Integrate the content filter with the extension-based verification.



- Parallel Sorting. Group strings by lengths using existing parallel algorithm.
- Parallel Building Indexes. Parallel building indexes for each group.
- Parallel Joins. Parallel perform similarity joins on each groups.

 Motivation
 Evaluating Pruning Techniques

 Dur Approach
 Evaluating Parallelism

 Experiment
 Evaluating Scalability

Experiment Setup

Table: Datasets

Datasets	cardinality	average len	max len	min len
GeoNames	400,000	11.106	1	60
GeoNames Query	100,000	10.7	2	43
Reads	750,000	101.388	86	106
Reads Query	100,000	101.2	88	116

MotivationEvaluating Pruning TechniquesOur ApproachEvaluating ParallelismExperimentEvaluating Scalability

Experiment Setup



Figure: Length Distribution.

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □



Evaluating Pruning Techniques



Figure: Evaluating pruning techniques for similarity joins(8 threads).



Evaluating Pruning Techniques



Figure: Evaluating pruning techniques for similarity search(8 threads).

Motivation Evaluating Pruning Technique Our Approach Evaluating Parallelism Experiment Evaluating Scalability

Evaluating Parallelism



Figure: Evaluating running time of similarity join by varying number of threads.

Motivation Evaluating Pruning Technique Our Approach Evaluating Parallelism Experiment Evaluating Scalability

Evaluating Speedup



Figure: Evaluating speedup of similarity join.

<ロ> <同> <同> <同> <同> <同> <同> <同> <同> <同</p>

 Motivation
 Evaluating Pruning Technique

 Our Approach
 Evaluating Parallelism

 Experiment
 Evaluating Scalability

Evaluating Parallelism



Figure: Evaluating running time of similarity search by varying number of threads.

イロト イポト イヨト イヨ

31= 990

Motivation Evaluating Pruning Technique Our Approach Evaluating Parallelism Experiment Evaluating Scalability

Evaluating Speedup



Figure: Evaluating speedup of similarity search.

◆□▶ ◆□▶ ◆□▶ ◆□▶ ●□□ のQ@

 Motivation
 Evaluating Pruning Techniques

 Our Approach
 Evaluating Parallelism

 Experiment
 Evaluating Scalability

Evaluating Scalability



Figure: Evaluating the scalability of the similarity join algorithm(8 threads).

 Motivation
 Evaluating Pruning Techniques

 Our Approach
 Evaluating Parallelism

 Experiment
 Evaluating Scalability

Evaluating Scalability



Figure: Evaluating the scalability of the similarity search algorithm(8 threads).

About our team I

- We are from Tsinghua University, Beijing, China.
- Yu Jiang, Jiannan Wang, Guoliang Li, Jianhua Feng and Dong Deng.

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □

Appendix

Our Team

About our team II



Dong Deng Parallel PassJoin

Thank You Q & A

http://dbgroup.cs.tsinghua.edu.cn/dd

Pass-Join: A Partition based Method for Similarity Joins. Guoliang Li, Dong Deng, Jiannan Wang, Jianhua Feng. VLDB 2012.