

Similarity Query Processing for Probabilistic Sets

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Motivation

- Evaluate similarity between uncertain sets
- Existing work
 - Huge model size
 - Significant similarity evaluation cost
- This paper
 - Comprehensive study for probabilistic set may have thousands of elements

Solution

- Similarities based on dynamic programming
 - Expected Similarity (ES)
 - Confidence-based Similarity (CS)
- Exact query processing based on pruning
 - Individual pruning
 - Batch pruning
- Approximate query processing based on sampling

Agenda

- Introduction
- Related work
- Problem definition and data normalization
- Exact similarity computation
- Pruning techniques
- Approximate solution
- Experiments

Introduction

- Applications
 - Personalization systems
 - Multi-label classification
- Contribution
 - Handle large p-sets efficiently
 - Similarity measure based on dynamic programming
 - Pruning techniques and approximate methods
 - Experiments upon synthetic and real datasets

Related work

- Uncertain Data Management
 - Information extraction and integration, multimedia retrieval, optical character recognition
 - MayBMS, MystiQ, Trio
- Similarity Search
 - Top-k, k-NN, reverse k-NN, range queries
- Similarity Join
 - Batch similarity queries

Related work

- Efficient processing of probabilistic set-containment queries on uncertain set-valued data.[*Inf. Sci*]
 - Same
 - Probabilistic set model, one of the similarity measure
 - Different
 - Pruning methods, approximate methods
- Probabilistic string similarity joins.[*SIGMOD 2010*]
 - Different
 - Non-neglectible correlations
 - Involving aggregated probabilities

Related work

- Set similarity join on probabilistic data.[*VLDB 2010*]

Model	Expressive Power	Exact Similarity Computation	Upper Bound Computation
Set-level [27]	Most general	$O(N^2)$	$O(N)$
Element-level [27]	Can model exclusion	$\Omega(2^n)$	$O(n^2)$ (online) or $O(n)$ (offline)
Our p-set model	A special case of Element-level model	$O(n^3)$	$O(n)$

– Models and Similarity Evaluation

- Set-level
- Element-level

– Pruning Rules

- Jaccard Distance pruning
- Probability upper bound pruning

Problem definition and data normalization

- Probabilistic set model

$$\mathcal{A} = \{a_i : p_{a_i} | a_i \in \mathcal{D}, \forall i \in [1, n]\}$$

- Possible world semantics

$$w(\mathcal{A}) \qquad \Pr[w] = \prod_{t \in w} p_t \prod_{t \notin w} (1 - p_t)$$

$$\mathcal{W}(\mathcal{A}, \mathcal{B}) = \mathcal{W}(\mathcal{A}) \times \mathcal{W}(\mathcal{B}) \qquad (w_a, w_b) \in \mathcal{W}(\mathcal{A}, \mathcal{B}) \text{ is } \Pr[w_a] \cdot \Pr[w_b].$$

- Jaccard coefficient

$$jac(X, Y) = \frac{|X \cap Y|}{|X \cup Y|}$$

Problem definition and data normalization

- Example
 - P-sets

\mathcal{A}	\mathcal{B}
$\{1 : 0.7, 2 : 1.0\}$	$\{1 : 1.0, 2 : 0.5, 3 : 0.8\}$

- All the joint possible worlds

w_a	w_b	$\Pr[(w_a, w_b)]$	Jaccard
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}\}$	0.03	0
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}\}$	0.03	0.5
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.12	0
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.12	0.333
$\{2^{\mathcal{A}}, 1^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}\}$	0.07	0.5
$\{2^{\mathcal{A}}, 1^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}\}$	0.07	1
$\{2^{\mathcal{A}}, 1^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.28	0.333
$\{2^{\mathcal{A}}, 1^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.28	0.666

Problem definition and data normalization

- Expected Similarity (ES)

$$\begin{aligned} ES(\mathcal{A}, \mathcal{B}) &= \sum_{(w_a, w_b) \in \mathcal{W}(\mathcal{A}, \mathcal{B})} sim(w_a, w_b) \cdot \Pr[(w_a, w_b)] \\ &= \sum_{w_a \in \mathcal{W}(\mathcal{A}) \wedge w_b \in \mathcal{W}(\mathcal{B})} sim(w_a, w_b) \cdot \Pr[w_a] \cdot \Pr[w_b] \end{aligned}$$

- Confidence-based Similarity (CS)

$$CS(\mathcal{A}, \mathcal{B}, minconf) = \max\{x \mid CPr(x, \mathcal{A}, \mathcal{B}) \geq minconf\}$$

– conditioned cumulative probability $CPr(x, \mathcal{A}, \mathcal{B})$

$$CPr(x, \mathcal{A}, \mathcal{B}) = \sum_{(w_a, w_b) \in \mathcal{W}(\mathcal{A}, \mathcal{B}) \wedge sim(w_a, w_b) \geq x} \Pr[(w_a, w_b)]$$

Problem definition and data normalization

- Example

\mathcal{A}	\mathcal{B}
$\{1 : 0.7, 2 : 1.0\}$	$\{1 : 1.0, 2 : 0.5, 3 : 0.8\}$

w_a	w_b	$\Pr[(w_a, w_b)]$	Jaccard
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}\}$	0.03	0
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}\}$	0.03	0.5
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.12	0
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.12	0.333
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$\{2^{\mathcal{A}}, 1^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.28	0.666

	Jaccard
$ES(\mathcal{A}, \mathcal{B})$	0.44
$CS(\mathcal{A}, \mathcal{B}, minconf = 0.3)$	0.666
$CS(\mathcal{A}, \mathcal{B}, minconf = 0.5)$	0.333

Problem definition and data normalization

- Normalization of two p-sets

$$\mathcal{A} = \{ c_1 : p_{c_1}^{\mathcal{A}}, \dots, c_k : p_{c_k}^{\mathcal{A}}, d_1 : p_{d_1}, \dots, d_{n-k} : p_{d_{n-k}} \}$$

$$\mathcal{B} = \{ c_1 : p_{c_1}^{\mathcal{B}}, \dots, c_k : p_{c_k}^{\mathcal{B}}, d_{n-k+1} : p_{d_{n-k+1}}, \dots, d_{n+m-2k} : p_{d_{n+m-2k}} \}$$

\mathcal{A}	\mathcal{B}
$\{1 : 0.7, 2 : 1.0\}$	$\{1 : 1.0, 2 : 0.5, 3 : 0.8\}$

- Size and expected size

w_a	w_b	$\Pr[(w_a, w_b)]$	Jaccard
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}\}$	0.03	0
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}\}$	0.03	0.5
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.12	0
$\{2^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.12	0.333
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$\{2^{\mathcal{A}}, 1^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.28	0.333
$\{2^{\mathcal{A}}, 1^{\mathcal{A}}\}$	$\{1^{\mathcal{B}}, 2^{\mathcal{B}}, 3^{\mathcal{B}}\}$	0.28	0.666

Exact similarity computation

- Equivalent classes

$$H[i, j] = \sum_{(w_a, w_b) \in \mathcal{W}(\mathcal{A}, \mathcal{B}) \wedge |w_a \cap w_b| = i \wedge |w_a \cup w_b| = j} \Pr[w_a] \cdot \Pr[w_b]$$

- Example

w_a	w_b	$\Pr[(w_a, w_b)]$	i	j	Jaccard
$\{2^A\}$	$\{1^B\}$	0.03	0	2	0
$\{2^A\}$	$\{1^B, 3^B\}$	0.12	0	3	0
$\{2^A\}$	$\{1^B, 2^B, 3^B\}$	0.12	1	3	0.333
$\{2^A, 1^A\}$	$\{1^B, 3^B\}$	0.28	1	3	0.333
$\{2^A\}$	$\{1^B, 2^B\}$	0.03	1	2	0.5
$\{2^A, 1^A\}$	$\{1^B\}$	0.07	1	2	0.5
$\{2^A, 1^A\}$	$\{1^B, 2^B, 3^B\}$	0.28	2	3	0.666
$\{2^A, 1^A\}$	$\{1^B, 2^B\}$	0.07	2	2	1

$H[i, j]$

	$j = 0$	$j = 1$	$j = 2$	$j = 3$
$i = 0$	0	0	0.03	0.12
$i = 1$		0	0.1	0.4
$i = 2$			0.07	0.28

Exact similarity computation

- Calculate ES

$$ES = \sum_{i=1}^k \sum_{j=i}^{m+n-k} H[i, j] \cdot (i/j)$$

- Calculate CS

Algorithm 1: Calculate CS from $H[i, j]$

Input: $H[i, j]$, $minconf$

Data: $heap$ is a max-heap on the similarity values.

```
1 for  $i = 1$  to  $k$  do  $heap.push(1.0, i, i)$ ;  
2  $CPr \leftarrow 0$ ;  $sim \leftarrow 0$ ;  
3 while  $heap.empty = false$  do  
4    $(sim, i, j) \leftarrow heap.pop$ ;  
5    $CPr \leftarrow CPr + H[i, j]$ ;  
6   if  $CPr \geq minconf$  then break;  
7   if  $j < m + n - k$  then  $heap.push(\frac{i}{j+1}, i, j + 1)$ ;  
8 return  $sim$ 
```

$H[i, j]$

	$j = 0$	$j = 1$	$j = 2$	$j = 3$
$i = 0$	0	0	0.03	0.12
$i = 1$		0	0.1	0.4
$i = 2$			0.07	0.28

Exact similarity computation

- Computing H
 - Common element

$$\begin{aligned} H^l[i, j] = & H^{l-1}[i, j](1 - p_l^A)(1 - p_l^B) \\ & + H^{l-1}[i, j-1](p_l^A(1 - p_l^B) + (1 - p_l^A)p_l^B) \\ & + H^{l-1}[i-1, j-1]p_l^A p_l^B \end{aligned}$$

- Distinct element

$$H^l[i, j] = H^{l-1}[i, j](1 - p_l) + H^{l-1}[i, j-1]p_l$$

- Time complexity $O(n^3)$
- Space complexity $O(n^2)$

$H[i, j]$

	$j = 0$	$j = 1$	$j = 2$	$j = 3$
$i = 0$	0	0	0.03	0.12
$i = 1$		0	0.1	0.4
$i = 2$			0.07	0.28

Pruning Techniques

Algorithm 2: Answer Queries with Pruning (\mathcal{Q} , $\{\mathcal{O}_i\}$, τ , $minconf$)

```

1  $C \leftarrow$  candidates that survive the batch pruning (c.f., Sec. V-D);
2 foreach  $p$ -set in  $C$  do
3    $pruned \leftarrow \text{false}$ ;
4   if the query type is ESQ then
5      $ub \leftarrow \text{calcESUpperBound}(\mathcal{Q}, \mathcal{O}_i)$  (c.f., Sec. V-B);
6     if  $ub < \tau$  then  $pruned \leftarrow \text{true}$ 
7   if the query type is CSQ then
8      $ub \leftarrow \text{calcCSUpperBound}(\mathcal{Q}, \mathcal{O}_i, \tau)$  (c.f., Sec. V-C);
9     if  $ub < minconf$  then  $pruned \leftarrow \text{true}$ 
10  if  $pruned = \text{false}$  then
11     $sim \leftarrow$  the similarity value between  $\mathcal{Q}$  and  $\mathcal{O}_i$ ;
12    if  $sim \geq \tau$  then
13      output  $\mathcal{O}_i$ ;

```

$$\mathbf{E}[|\mathcal{A}|], \text{ is } \sum_{w \in \mathcal{W}(\mathcal{A})} |w| \cdot \mathbf{Pr}[w] = \sum_{l=1}^n p_l^{\mathcal{A}}$$

$$\mathbf{E}[|\mathcal{A} \cap \mathcal{B}|] \text{ is } \sum_{(w_a, w_b) \in \mathcal{W}(\mathcal{A}, \mathcal{B})} |w_a \cap w_b| \cdot \mathbf{Pr}[(w_a, w_b)] = \sum_{l=1}^k p_l^{\mathcal{A}} \cdot p_l^{\mathcal{B}}$$

$$\mathbf{E}[|\mathcal{A} \cup \mathcal{B}|] \text{ is } \mathbf{E}[|\mathcal{A}|] + \mathbf{E}[|\mathcal{B}|] - \mathbf{E}[|\mathcal{A} \cap \mathcal{B}|] = \sum_{l=1}^k (p_l^{\mathcal{A}} + p_l^{\mathcal{B}} - p_l^{\mathcal{A}} \cdot p_l^{\mathcal{B}}) + \sum_{l=k+1}^{n+m-k} p_l$$

Pruning Techniques

- Pruning Rule for ESQ

$$\mathbf{E}[X/Y] < UB_1(\mathbf{E}[X], \mathbf{E}[Y])$$

$$UB_1(u, v) = \min_{\exp(-u/3) \leq \epsilon \leq 1} \left(2\epsilon + \frac{u + \sqrt{-3u \ln \epsilon}}{v - \sqrt{-2v \ln \epsilon}} \right)$$

$$UB_1(\mathbf{E}[|Q \cap \mathcal{O}|], \mathbf{E}[|Q \cup \mathcal{O}|]) \leq \tau$$

- Pruning Rule for CSQ

$$\Pr[X \geq \alpha Y] < UB_2(\mathbf{E}[X], \mathbf{E}[Y], \alpha)$$

$$UB_2(u, v, \alpha) = \min_{u \leq \xi \leq \min(\alpha v, 2u)} \left(e^{\frac{-(\alpha v - \xi)^2}{2\alpha^2 v}} + e^{\frac{-(\xi - u)^2}{3u}} \right)$$

$$\mathbf{E}[|Q \cap \mathcal{O}|] \leq \tau \cdot \mathbf{E}[|Q \cup \mathcal{O}|] \quad UB_2(\mathbf{E}[|Q \cap \mathcal{O}|], \mathbf{E}[|Q \cup \mathcal{O}|], \tau) \leq minconf$$

Pruning Techniques

- Batch Pruning
 - Discard many p-sets in the database without even evaluating their similarity upper bounds
 - Index all the p-sets in the database by their expected sizes
 - Compute a lower bound S_L and an upper bound S_U of the expected size for the appropriate query type
 - Only consider p-sets in the database whose expected sizes fall within $[S_L, S_U]$

Pruning Techniques

- Batch Pruning

- How to decide S_L and S_U

$$\mathbf{E}[|Q \cap \mathcal{O}|] \leq \min(\mathbf{E}[|Q|], \mathbf{E}[|\mathcal{O}|])$$

- Batch Pruning for ESQ

$$\mathbf{E}[|Q \cup \mathcal{O}|] \geq \max(\mathbf{E}[|Q|], \mathbf{E}[|\mathcal{O}|])$$

$$x + \sqrt{-3x \ln \epsilon^*} = (\tau - 2\epsilon^*)(\mathbf{E}[|Q|] - \sqrt{-2\mathbf{E}[|Q|] \ln \epsilon^*})$$

$$x - \sqrt{-2x \ln \epsilon^*} = \left(\mathbf{E}[|Q|] + \sqrt{-3\mathbf{E}[|Q|] \ln \epsilon^*} \right) / (\tau - 2\epsilon^*)$$

- Batch Pruning for CSQ

$$\exp\left(\frac{-(\xi_1^* - x)^2}{3x}\right) = \text{minconf}/2$$

$$\exp\left(\frac{-(\tau \cdot x - \xi_2^*)^2}{2\tau^2 \cdot x}\right) = \text{minconf}/2$$

Approximate solution

- Sampling-based method
 - Approximate algorithm for ES

$$\lceil (\ln \frac{2}{\delta}) / (2\epsilon^2) \rceil \quad \Pr \left[\left| \widehat{ES} - ES \right| \leq \epsilon \right] \geq 1 - \delta$$

- Approximate algorithm for CS

$$G = 24 \cdot \lceil \ln \frac{1}{\delta} \rceil, M = \lceil 2\epsilon^{-2} \rceil \quad \Pr \left[CS^- \leq \widehat{CS} \leq CS^+ \right] \geq 1 - \delta$$

- $O(n)$

Experiments

- Implementation
 - Java
 - Intel Pentium IV 2.8GHz CPU
 - 4GB memory
- Synthetic datasets
 - SYN*a*-U
 - a uniform distribution within the range of $[v, 0.9]$ with a default v value of 0.2.
 - SYN*a*-G
 - a Gaussian distribution $N(u, o)$ capped to the range of $(0, 1]$. By default, $u = 0.8$ and $o = 0.2$.

Experiments

- Real-world datasets

Dataset	DB Size	p-set Min/Max/Avg Size
pDBLP	5,000	27 / 708 / 204.9
pDeli	44,876	50 / 293,214 / 453.2

- pDBLP

- a fairly simple yet effective method based on topical terms used in authors' DBLP entries

- pDeli

- the social bookmarking dataset which was crawled from the Del.icio.us web site during 2006 and 2007

- Sigmoid function

$$p(e) = \frac{2}{1 + \exp(-c(e))} - 1$$

Experiments

- Default parameters

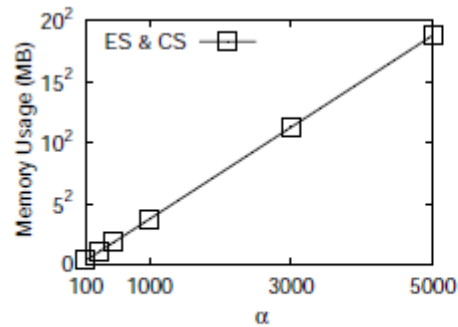
are: $minconf = 0.5$ (for CS), $\tau = 0.5$, $\alpha = 1000$, $\gamma = 10\%$,
 $\epsilon = 0.06$, $\delta = 0.06$, $v = 0.2$, $\sigma = 0.2$, and $\mu = 0.8$.

- Measures

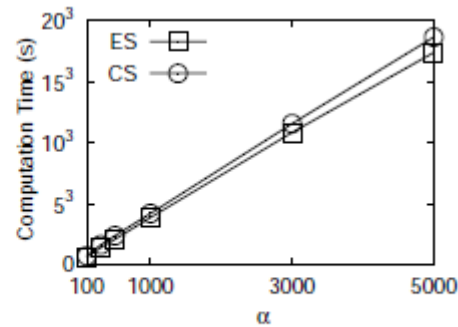
- Memory Usage
- Computation Time
- Query Time, Pruning time
- Candidate size, result size
- Pruning rate
- Average precision

Experiments

- Computing Similarities Exactly



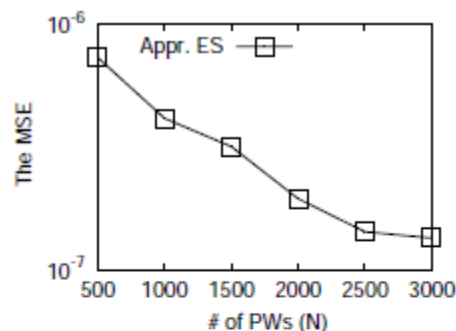
(a) Space consumption



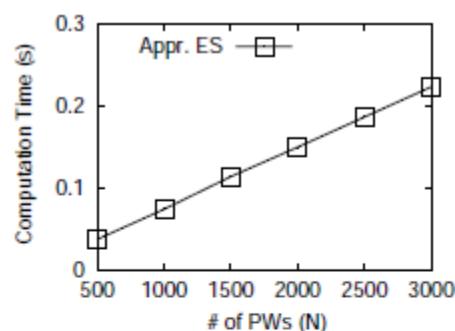
(b) Computation time

Experiments

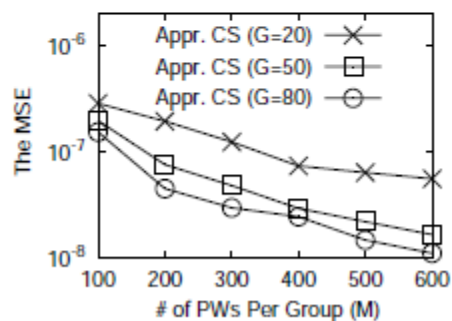
- Computing Similarities Approximately



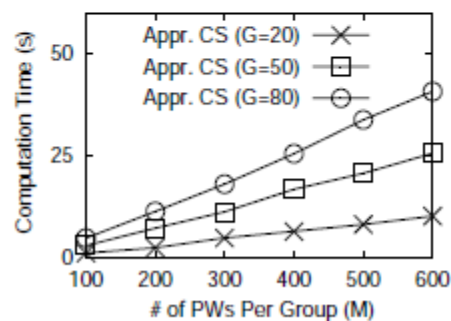
(a) ES, MSE



(b) ES, Computation time



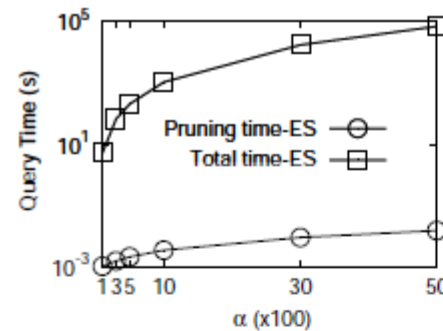
(c) CS, MSE



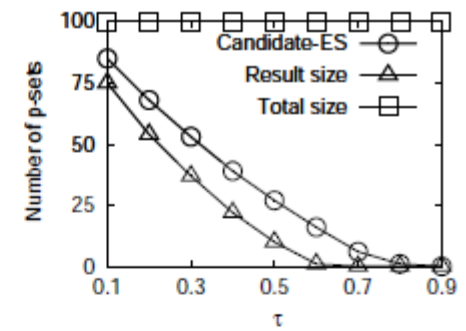
(d) CS, Computation time

Experiments

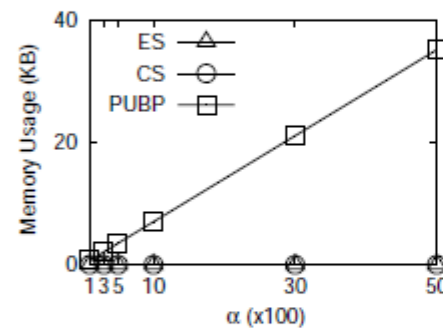
- Evaluating Pruning Efficiency on SYN



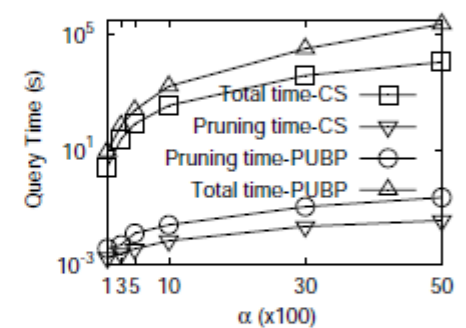
(a) *ESQ*, Query Time



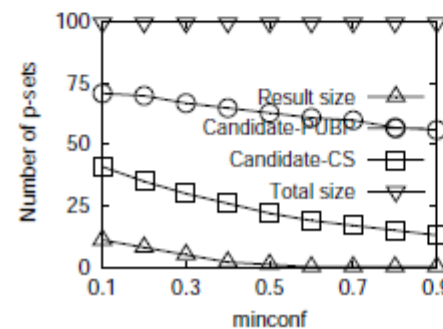
(b) *ESQ*, Candidate Size



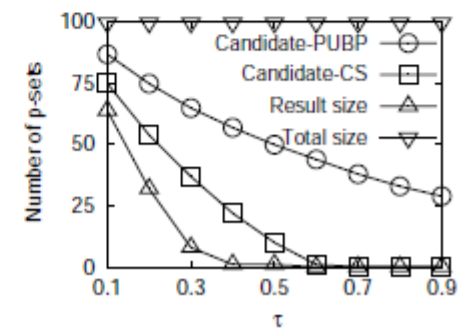
(c) *CSQ*, Memory Usage



(d) *CSQ*, Query Time



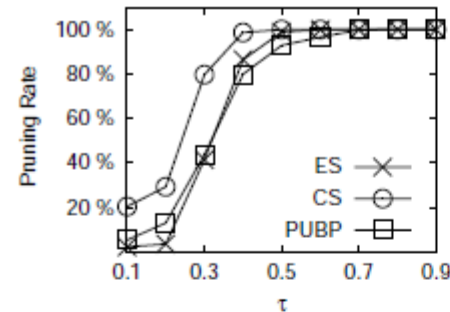
(e) *CSQ*, Candidate Size



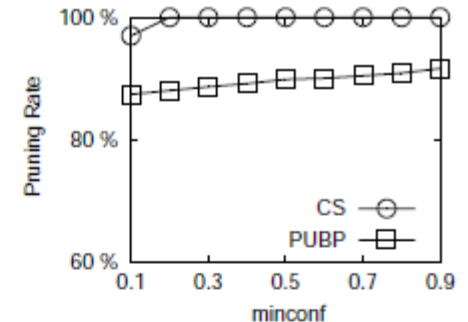
(f) *CSQ*, Candidate Size

Experiments

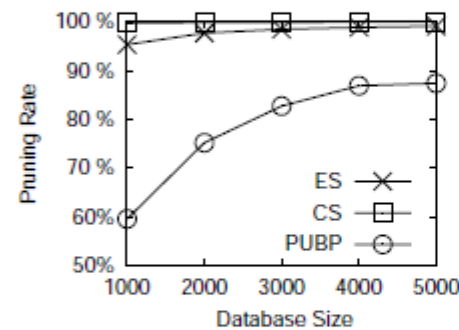
- Performance on the pDBLP Dataset



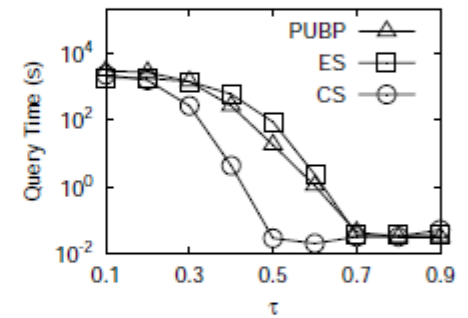
(a) Pruning Rate



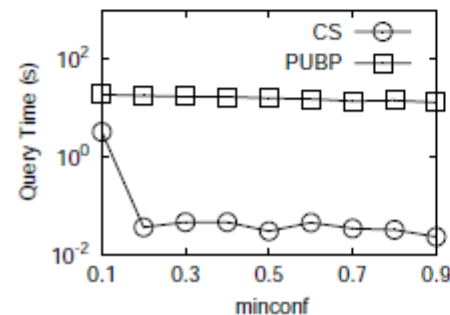
(b) Pruning Rate



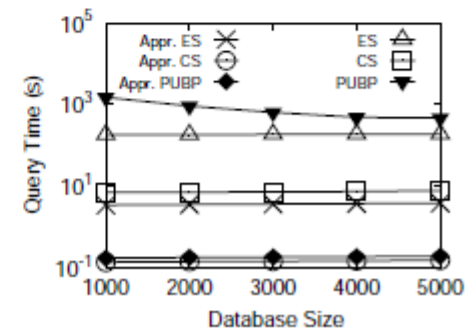
(c) Pruning Rate



(d) Query Time



(e) Query Time

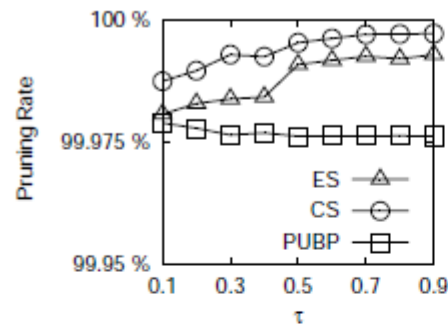


(f) Query Time

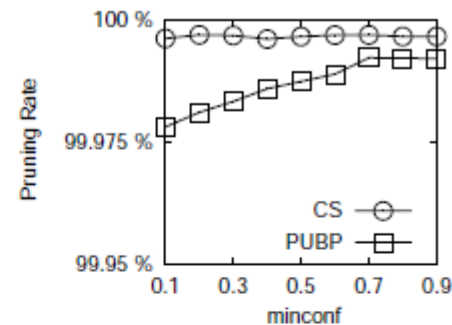
Experiments

AP@ k	5	10	15	20	25	30
ES	0.7000	0.6675	0.6250	0.5875	0.5825	0.5500
CS	0.7000	0.6785	0.6280	0.5825	0.575	0.5500

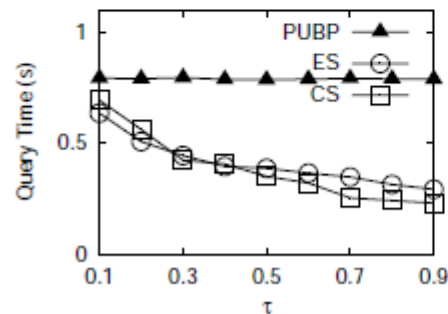
- Performance on the pDeli Dataset



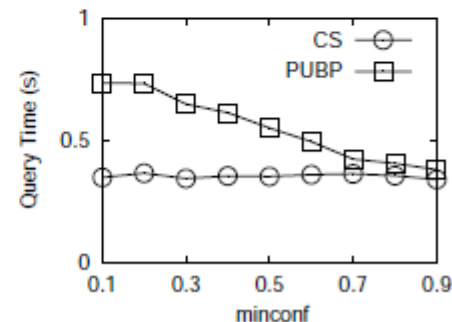
(a) Pruning Rate



(b) Pruning Rate



(c) Query Time



(d) Query Time

Thank You!