

Basic principles of algorithmic graph mining Lecture 2 : Computing basic graph statistics

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Estonian Summer School on Computer and System Science 2016
Aug 22–24, 2016

course agenda

- introduction to graph mining
- computing basic graph statistics
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- spectral graph analysis
- additional topics and applications

algorithmic tools

efficiency considerations

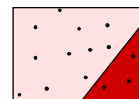
- data in the web and social-media are typically of extremely large scale (easily reach to billions)
- how to compute simple graph statistics?
- even quadratic algorithms are not feasible in practice

hashing and sketching

- probabilistic / approximate methods
- sketching : create sketches that summarize the data and allow to estimate simple statistics with small space
- hashing : hash objects in such a way that similar objects have larger probability of mapped to the same value than non-similar objects

estimator theorem

- consider a set of items U
- a fraction ρ
- estimate ρ by sampling



- how many samples N are needed?

$$N \geq \frac{4}{\epsilon^2 \rho} \log \frac{2}{\delta}$$

for an ϵ -approximation with probability at least $1 - \delta$

- notice: it does not depend on $|U|$ (!)

homework

use the Chernoff bound to derive the estimator theorem

computing statistics on data streams

- $X = (x_1, x_2, \dots, x_m)$ a sequence of elements
- each x_i is a member of the set $N = \{1, \dots, n\}$
- $m_i = |\{j : x_j = i\}|$ the number of occurrences of i

$$F_k = \sum_{i=1}^n m_i^k$$

computing statistics on data streams

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- F_0 is the number of distinct elements

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- F_1 is the length of the sequence

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$$F_k = \sum_{i=1}^n m_i^k$$

- F_0 is the number of distinct elements
- F_1 is the length of the sequence
- F_2 index of homogeneity, size of self-join, and other applications

computing statistics on data streams

- How to compute the frequency moments using less than $O(n \log m)$ space?
- **sketching**: create a sketch that takes much less space and gives an estimation of F_k

[Alon et al., 1999]

estimating the number of distinct values (F_0)

[Flajolet and Martin, 1985]

- consider a bit vector of length $O(\log n)$
- upon seen x_i , **set**:
 - the 1st bit with probability $1/2$
 - the 2nd bit with probability $1/4$
 - ...
 - the i -th bit with probability $1/2^i$
- **important**: bits are set deterministically for each x_i
- let R be the index of the largest bit set
- return $Y = 2^R$

estimating number of distinct values (F_0)

Theorem. For every $c > 2$, the algorithm computes a number Y using $O(\log n)$ memory bits, such that the probability that the ratio between Y and F_0 is not between $1/c$ and c is at most $2/c$

locality sensitive hashing

a family \mathcal{H} is called (R, cR, p_1, p_2) -sensitive if for any two objects p and q

- if $d(p, q) \leq R$, then $\Pr_{\mathcal{H}}[h(p) = h(q)] \geq p_1$
- if $d(p, q) \geq cR$, then $\Pr_{\mathcal{H}}[h(p) = h(q)] \leq p_2$

interesting case when $p_1 > p_2$

locality sensitive hashing: example

- objects in a Hamming space $\{0, 1\}^d$ – binary vectors
- $\mathcal{H} : \{0, 1\}^d \rightarrow \{0, 1\}$ sample the i bit:
- $\mathcal{H} = \{h(x) = x_i \mid i = 1, \dots, d\}$
- for two vectors x and y with distance r , it is

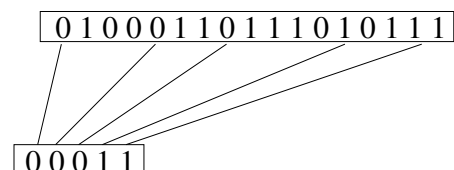
$$\Pr_{\mathcal{H}}[h(x) = h(y)] = 1 - \frac{r}{d}$$

- thus $p_1 = 1 - \frac{R}{d}$ and $p_2 = 1 - \frac{cR}{d}$
- gap between p_1 and p_2 too small
- probability amplification

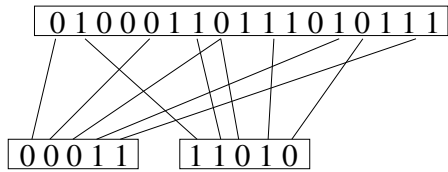
locality sensitive hashing: Hamming distance

0 1 0 0 0 1 1 0 1 1 1 0 1 0 1 1 1

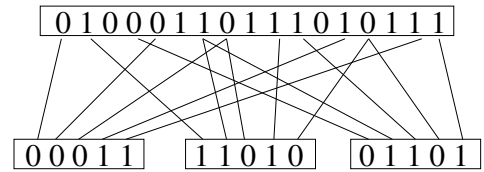
locality sensitive hashing: Hamming distance



locality sensitive hashing: Hamming distance



locality sensitive hashing: Hamming distance

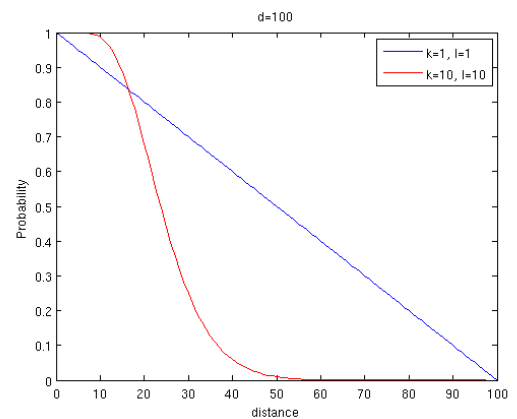


locality sensitive hashing: Hamming distance

Probability of collision

$$\Pr[h(x) = h(y)] = 1 - (1 - (1 - \frac{r}{d})^k)^l$$

locality sensitive hashing: Hamming distance



homework

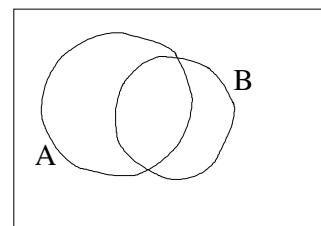
how to apply the locality sensitive hashing for vectors of integers, not just binary vectors?

vectors $\mathbf{x} = \{x_1, \dots, x_d\}$

L_1 distance $\|\mathbf{x} - \mathbf{y}\|_1 = \sum_{i=1}^d |x_i - y_i|$

Jaccard coefficient

- for two sets $A, B \subseteq U$ $J(A, B) = \frac{|A \cap B|}{|A \cup B|}$
- measure of similarity of the sets



- can we design a locality sensitive hashing family for Jaccard?

min-wise independent permutations

- $\pi : U \rightarrow U$ a random permutation of U
- $h(A) = \min\{\pi(x) \mid x \in A\}$

min-wise independent permutations

- $\pi : U \rightarrow U$ a random permutation of U
- $h(A) = \min\{\pi(x) \mid x \in A\}$
- then

$$\Pr[h(A) = h(B)] = J(A, B) = \frac{|A \cap B|}{|A \cup B|}$$

- amplify the probability as before:
 - repeat many times,
 - concatenate into blocks
 - consider objects similar if they collide in at least one block

homework

show that for $h(A) = \min\{\pi(x) \mid x \in A\}$, with π a random permutation, it is

$$\Pr[h(A) = h(B)] = J(A, B) = \frac{|A \cap B|}{|A \cup B|}$$

homework

design a locality-sensitive hashing scheme for vectors according to the cosine similarity measure

vectors $\mathbf{x} = \{x_1, \dots, x_d\}$

distance $1 - \cos(\mathbf{x}, \mathbf{y}) = 1 - \frac{\mathbf{x} \cdot \mathbf{y}}{\|\mathbf{x}\|_2 \|\mathbf{y}\|_2}$

applications of the algorithmic tools to real scenarios

clustering coefficient and triangles

clustering coefficient

$$C = \frac{3 \times \text{number of triangles in the network}}{\text{number of connected triples of vertices}}$$

- how to compute it?
- how to compute the number of triangles in a graph?
- assume that the graph is very large, stored in disk

[Buriol et al., 2006]

- count triangles when graph is seen as a data stream
- two models:
 - edges are stored in any order
 - edges in order : all edges incident to one vertex are stored sequentially

counting triangles

- brute-force algorithm is checking every triple of vertices
- obtain an approximation by sampling triples



sampling algorithm for counting triangles



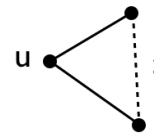
- how many samples are required?
- let T be the set of all triples and T_i the set of triples that have i edges, $i = 0, 1, 2, 3$
- by the **estimator theorem**, to get an ϵ -approximation, with probability $1 - \delta$, the number of samples should be

$$N \geq O\left(\frac{|T|}{|T_3|} \frac{1}{\epsilon^2} \log \frac{1}{\delta}\right)$$

- but $|T|$ can be very large compared to $|T_3|$

counting triangles

- **incidence model** : all edges incident to each vertex appear **in order** in the stream
- sample connected triples



sampling algorithm for counting triangles

- incidence model
- consider sample space $\mathcal{S} = \{b-a-c \mid (a, b), (a, c) \in E\}$
- $|\mathcal{S}| = \sum_i d_i(d_i - 1)/2$

- 1: sample $X \subseteq \mathcal{S}$ (paths $b-a-c$)
- 2: estimate fraction of X for which edge (b, c) is present
- 3: scale by $|\mathcal{S}|$

- gives (ϵ, δ) approximation

counting triangles — incidence stream model

SAMPLETRIANGLE [Buriol et al., 2006]

1st pass

count the number of paths of length 2 in the stream

2nd pass

uniformly choose one path (a, b, c)

3rd pass

if $((b, c) \in E)$ $\beta = 1$ else $\beta = 0$

return β

counting triangles — incidence stream model

SAMPLETRIANGLE [Buriol et al., 2006]

1st pass

count the number of paths of length 2 in the stream

2nd pass

uniformly choose one path (a, b, c)

3rd pass

if $((b, c) \in E) \beta = 1$ else $\beta = 0$

return

we have $E[\beta] = \frac{3|T_3|}{|T_2|+3|T_3|}$, with $|T_2| + 3|T_3| = \sum_u \frac{d_u(d_u-1)}{2}$, so

$$|T_3| = E[\beta] \sum_u \frac{d_u(d_u-1)}{6}$$

and space needed is $O((1 + \frac{|T_2|}{|T_3|}) \frac{1}{\epsilon^2} \log \frac{1}{\delta})$

properties of the sampling space

it should be possible to

- estimate the size of the sampling space
- sample an element uniformly at random

homework

- 1 compute triangles in 3 passes when edges appear in arbitrary order
- 2 compute triangles in 1 pass when edges appear in arbitrary order
- 3 compute triangles in 1 pass in the incidence model

triangle sparsifiers

[Tsourakakis et al., 2011]

- start with graph $G(V, E)$
- use sparsification parameter p
- pick a random subset E' of edges
each edge is selected with probability p
- $T'_3 = \#$ triangles on graph $G'(V, E')$
- return $T_3 = T'_3/p^3$

triangle sparsifiers

[Tsourakakis et al., 2011]

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- use sparsification parameter p
- pick a random subset E' of edges
each edge is selected with probability p
- $T'_3 = \#$ triangles on graph $G'(V, E')$
- return $T_3 = T'_3/p^3$
- T_3 is highly concentrated around the true number of triangles

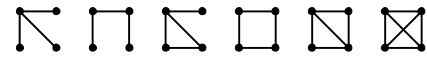
counting graph minors

counting other minors

- count all **minors** in a very large graphs
 - connected subgraphs
 - size 3 and 4
 - directed or undirected graphs
- why?**
- modeling networks, "**signature**" structures
e.g., copying model
- anomaly detection**, e.g., spam link farms
[Alon, 2007, Bordino et al., 2008]

counting minors in large graphs

- characterize** a graph by the distribution of its minors



all undirected minors of size 4



all directed minors of size 3

sampling algorithm for counting triangles

- incidence model
 - consider sample space $\mathcal{S} = \{b-a-c \mid (a,b), (a,c) \in E\}$
 - $|\mathcal{S}| = \sum_i d_i(d_i - 1)/2$
- sample $X \subseteq \mathcal{S}$ (paths $b-a-c$)
 - estimate fraction of X for which edge (b,c) is present
 - scale by $|\mathcal{S}|$
- gives (ϵ, δ) approximation

adapting the algorithm

sampling spaces:

- 3-node directed



- 4-node undirected



are the sampling space properties satisfied?

datasets

graph class	type	# instances
synthetic	un/directed	39
wikipedia	un/directed	7
webgraphs	un/directed	5
cellular	directed	43
citation	directed	3
food webs	directed	6
word adjacency	directed	4
author collaboration	undirected	5
autonomous systems	undirected	12
protein interaction	undirected	3
US road	undirected	12

clustering of undirected graphs

assigned to	0	1	2	3	4	5	6
AS graph	12	0	0	0	0	0	0
collaboration	0	0	3	2	0	0	0
protein	1	0	0	1	0	0	1
road-graph	0	12	0	0	0	0	0
wikipedia	0	0	0	0	2	5	0
synthetic	11	0	0	0	0	0	28
webgraph	2	0	0	1	0	0	0

clustering of directed graphs

feature class	accuracy compared to ground truth
standard topological properties (81)	0.74%
minors of size 3	0.78%
minors of size 4	0.84%
minors of size 3 and 4	0.91%

local statistics

compute local statistics in large graphs

- **our goal:** compute triangle counts for all vertices
-
- **motivation**
 - motifs can be used to characterize network families [Alon, 2007, Bordino et al., 2008]
 - analysis of social or biological networks
 - thematic relationships in the web
 - web spam
- **applications:** spam detection and content quality analysis in social media

semi-streaming model

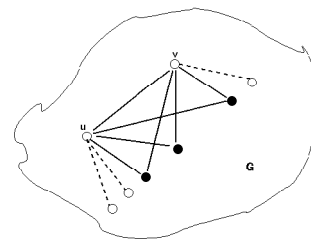
[Feigenbaum et al., 2004]

- data stream model (constant memory) **too restrictive**
- graph stored in **secondary memory** as adjacency or edge list
- ✗ no random access possible
- $O(N \log N)$ bits available in **main memory**
 - limited amount of information per vertex
 - ✗ not enough to store edges in main memory
- limited (constant or $O(\log N)$) number of passes
- compute counts for all vertices concurrently

two algorithms

- 1 **external memory**
 - keep a **counter** for **each vertex** (main memory)
 - keep a **counter** for **each edge** (secondary memory)
- 2 **main memory**
 - keep a **counter** for **each vertex**

number of triangles for edges and nodes



- neighbors: $N(u) = \{v : (u, v) \in E\}$
- degree: $d(u) = |N(u)|$
- edge triangles: $T_{uv} = |N(u) \cap N(v)|$
- vertex triangles: $T(u) = \frac{1}{2} \sum_{v \in N(u)} T_{uv}$

computing triangles : idea

- consider the Jaccard coefficient between two sets A and B :

$$J(A, B) = \frac{|A \cap B|}{|A \cup B|}$$

- if we knew $J(N(u), N(v)) = J$, then:

$$T_{uv} = |N(u) \cap N(v)| = \frac{J}{J+1} (|N(u)| + |N(v)|)$$

- and then:

$$T(u) = \frac{1}{2} \sum_{v \in N(u)} T_{uv}$$

computing triangles : idea

we want:

$$T_{uv} = |N(u) \cap N(v)| = \frac{J}{J+1} (|N(u)| + |N(v)|)$$

approximate

- m independent trials
- Z_{uv} : # times that $\min \pi(N(u)) = \min \pi(N(v))$

use the estimator:

$$\bar{T}_{uv} = \frac{Z_{uv}}{Z_{uv} + m} (|N(u)| + |N(v)|)$$

external-memory algorithm

- semi-stream model
- keep vertex min-hash values (in memory)
- keep edge counters (on disk)
- use edge counters to estimate number of triangles (and local clustering coefficient)

external-memory algorithm

```

1: Z = 0
2: for i: 1 ... m do {independent trials}
3:   for u : 1 ... |V| do {assign labels}
4:      $l_i(u) = \text{hash}_i(u)$  {Min-wise linear permutation}
5:   end for

```

external-memory algorithm

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6:   for u : 1 ... |V| do {compute fingerprints}
7:      $F_i(u) = \min_{v \in N(u)} l_i(u)$ 
8:   end for {1 scan of G}

```

external-memory algorithm

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8:   end for {1 scan of G}
9:   for u : 1 ... |V| do {update counters}
10:    for v in N(u) do
11:      if ( $F_i(u) = F_i(v)$ ) then {minima are equal}
12:         $Z_{uv} = Z_{uv} + 1$  { $Z_{uv}$ 's stored on disk}
13:      end if
14:    end for
15:   end for
16: end for

```

implementation

- $\text{hash}_i(x)$ is, e.g., a linear hash function ($a_i x + b_i \bmod p$)
- for every i , the $F_i(u)$'s can be kept in main memory
- the Z_{uv} 's must be stored on disk
 - for every i , updating Z_{uv} requires access to disk
 - computing counters most expensive operation

main-memory algorithm

- replace:
$$\bar{T}_{uv} = \frac{Z_{uv}}{Z_{uv} + m} (|N(u)| + |N(v)|)$$
- by the estimator for $|N(u) \cap N(v)|$:
$$\tilde{T}_{uv} = \frac{Z_{uv}}{\frac{2}{3}m} (N(u) + N(v))$$
- and estimator for $T(u)$:
$$\tilde{T}(u) = \frac{1}{3m} \sum_{v \in N(u)} Z_{uv} (N(u) + N(v)) = \frac{1}{3m} Z_u$$
- Z_u sums $d(u) + d(v)$ if $\min \pi(N(u)) = \min \pi(N(v))$
- only one counter per node

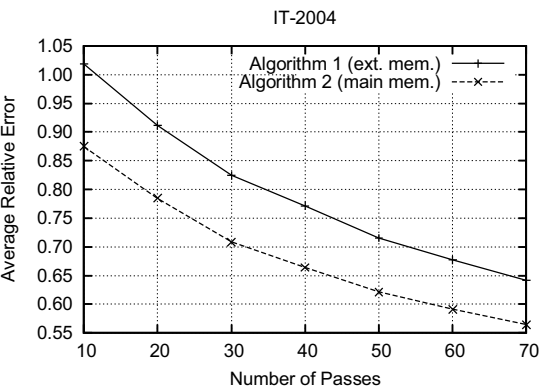
main-memory algorithm

```
1: Z = 0
2: for i: 1 ... m do {Independent trials}
3:   for u : 1 ... |V| do {Assign labels}
4:      $l_i(u) = \text{hash}_i(u)$ 
5:   end for
6:   for u : 1 ... |V| do
7:      $F_i(u) = \min_{v \in V(u)} l_i(u)$ 
8:   end for {1 scan of G}
9:   for u : 1 ... |V| do {Update counters}
10:    for  $v \in N(u)$  do
11:      if  $F_i(u) == F_i(v)$  then {Minima are equal}
12:         $Z_u = Z_u + d(u) + d(v)$  { $Z_u$ 's in main memory}
13:      end if
14:    end for
15:  end for
16: end for
```

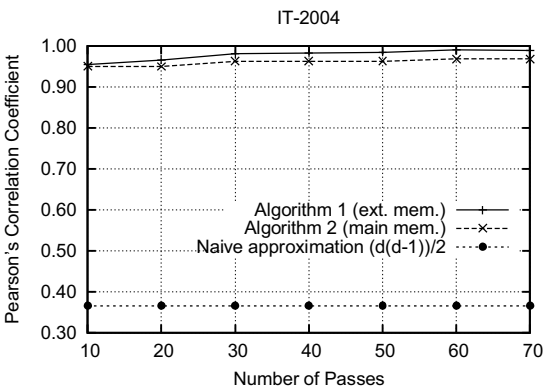
experimental results

Graph	Nodes	Edges	Algorithm 1 (ext. mem.)	Algorithm 2 (main mem.)
WB-2001	118M	1.7G	10 hr 20 min	3 hr 40 min
IT-2004	41M	2.1G	8 hr 20 min	5 hr 30 min
UK-2006	77M	5.3G	20 hr 30 min	13 hr 10 min

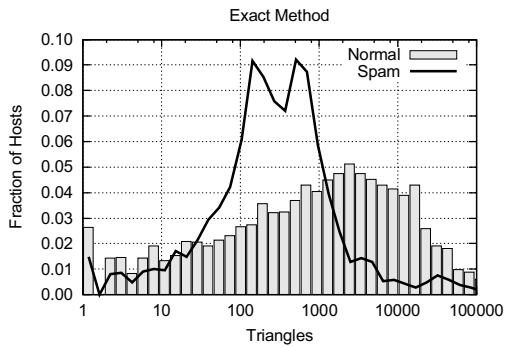
quality of approximation



quality of approximation

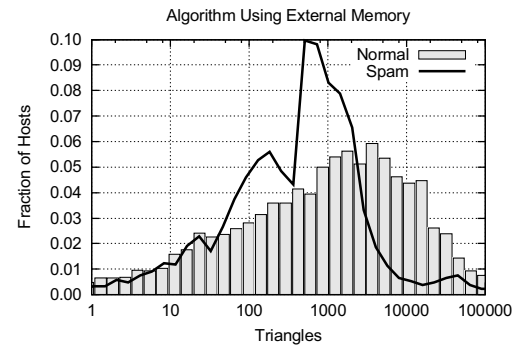


applications : spam detection



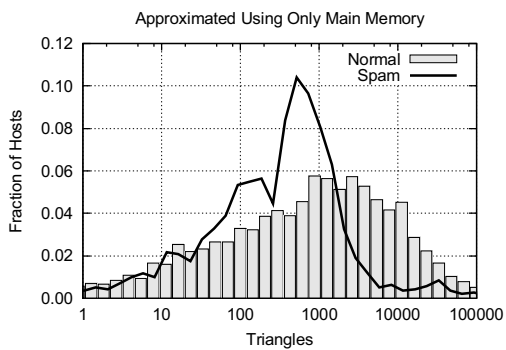
Separation of non-spam and spam hosts in the histogram of triangles

applications : spam detection



Separation of non-spam and spam hosts in the histogram of triangles

applications : spam detection



Separation of non-spam and spam hosts in the histogram of triangles

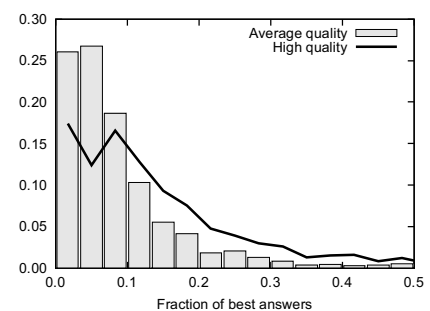
applications : spam detection

number of triangles feature is ranked 60-th out of 221 for spam detection

applications : content quality in yahoo! answers

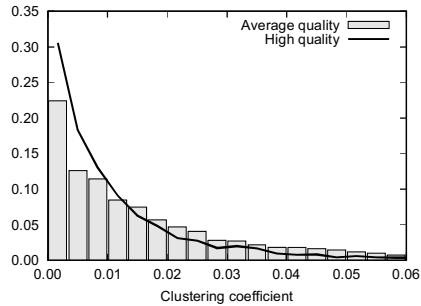
- **Yahoo! answers**, a question-answering portal
- consider the graph with edges (u, v) if user u has answered a question of user v
- consider **"high quality" users** those who have given a best answer to a random sample of questions
- **predict** high-quality users based on their local structure

applications : content quality in yahoo! answers



Separation of users who have provided questions/answers of high quality with users who have provided questions/answers of normal quality in terms of fraction of best answers

applications : content quality in yahoo! answers



Separation of users who have provided questions/answers of high quality with users who have provided questions/answers of normal

graph distance distributions

small-world phenomena

small worlds : graphs with short paths



- Stanley Milgram (1933-1984)
"The man who shocked the world"
- obedience to authority (1963)
- small-world experiment (1967)

Milgram's experiment

- 300 people (starting population) are asked to **dispatch a parcel** to a single individual (target)
- the target was a Boston stockbroker
- the starting population is selected as follows:
 - 100 were random **Boston inhabitants** (group A)
 - 100 were random **Nebraska stockbrokers** (group B)
 - 100 were random **Nebraska inhabitants** (group C)

Milgram's experiment

- **rules of the game** :
- parcels could be directly sent only to someone the sender **knows personally**
- 453 **intermediaries** happened to be involved in the experiments (besides the starting population and the target)

Milgram's experiment

questions Milgram wanted to answer:

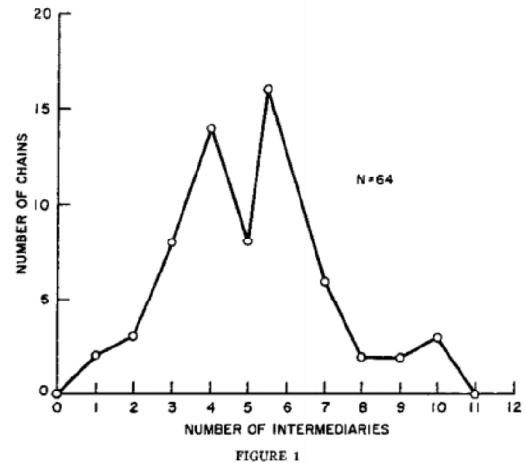
1. how many parcels will reach the target?
2. what is the distribution of the number of hops required to reach the target?
3. is this distribution different for the three starting subpopulations?

Milgram's experiment

answers to the questions

1. how many parcels will reach the target?
29%
2. what is the distribution of the number of hops required to reach the target?
average was 5.2
3. is this distribution different for the three starting subpopulations?
YES: average for groups A/B/C was 4.6/5.4/5.7

chain lengths



measuring what?

but what did Milgram's experiment reveal, after all?

1. the world is small
2. that people are able to exploit this smallness

graph distance distribution

- obtain information about a large graph, i.e., social network
- macroscopic level
- distance distribution
 - mean distance
 - median distance
 - diameter
 - effective diameter
 - ...

graph distance distribution

- given a graph, $d(x, y)$ is the length of the shortest path from x to y ∞ if one cannot go from x to y
- for undirected graphs, $d(x, y) = d(y, x)$
- for every t , count the number of pairs (x, y) such that $d(x, y) = t$
- the fraction of pairs at distance t is a distribution

exact computation

how can one compute the distance distribution?

- weighted graphs: Dijkstra (single-source: $O(m \log n)$),
- Floyd-Warshall (all-pairs: $O(n^3)$)
- in the unweighted case:
 - a single BFS solves the single-source version of the problem: $O(m)$
 - if we repeat it from every source: $O(nm)$

sampling pairs

- sample at random pairs of nodes (x, y)
- compute $d(x, y)$ with a BFS from x
- (possibly: reject the pair if $d(x, y)$

sampling pairs

- for every t , the fraction of sampled pairs that were found at distance t are an estimator of the value of the probability mass function
- takes a BFS for every pair — $O(m)$

sampling sources

- sample at random a source t
- compute a full BFS from t

sampling sources

- it is an unbiased estimator only for undirected and connected graphs
- uses anyway BFS...
 - ...not cache friendly
 - ... not compression friendly

idea : diffusion

[Palmer et al., 2002]

- let $B_t(x)$ be the ball of radius t around x (the set of nodes at distance $\leq t$ from x)
- clearly $B_0(x) = \{x\}$
- moreover $B_{t+1}(x) = \bigcup_{(x,y)} B_t(y) \cup \{x\}$
- so computing B_{t+1} from B_t just takes a single (sequential) scan of the graph

easy but costly

- every set requires $O(n)$ bits, hence $O(n^2)$ bits overall
- easy but costly
- too many!
- what about using approximated sets?
- we need probabilistic counters, with just two primitives: add and size
- very small!

estimating the number of distinct values (F_0)

- [Flajolet and Martin, 1985]
- consider a bit vector of length $O(\log n)$
- upon seen x_i , **set**:
 - the 1st bit with probability $1/2$
 - the 2nd bit with probability $1/4$
 - ...
 - the i -th bit with probability $1/2^i$
- **important**: bits are set deterministically for each x_i
- let R be the index of the largest bit set
- return $Y = 2^R$

ANF

- probabilistic counter for approximating the number of distinct values [Flajolet and Martin, 1985]
- ANF algorithm [Palmer et al., 2002] uses the original probabilist counters
- HyperANF algorithm [Boldi et al., 2011] uses HyperLogLog counters [Flajolet et al., 2007]

HyperANF

- HyperLogLog counter [Flajolet et al., 2007]
- with 40 bits you can count up to 4 billion with a standard deviation of 6%
- remember: one set per node

implementation tricks

[Boldi et al., 2011]

- use broad-word programming
- systolic computation for on-demand updates of counters
- exploit micro-parallelization of multicore architectures

performance

- HADI, a Hadoop-conscious implementation of ANF [Kang et al., 2011]
- takes 30 minutes on a 200K-node graph (on one of the 50 world largest supercomputers)
- HyperANF does the same in 2.25min on a workstation (20 min on a laptop).

experiments on facebook

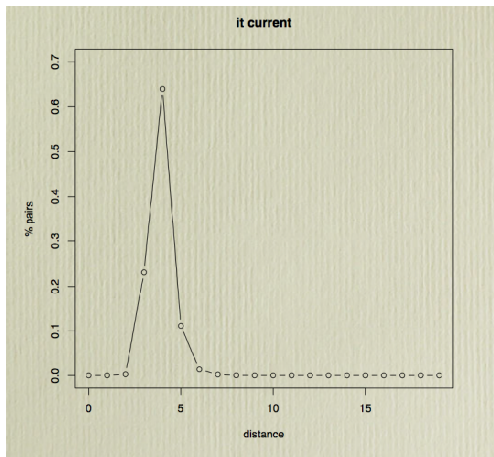
[Backstrom et al., 2011]

considered only active users

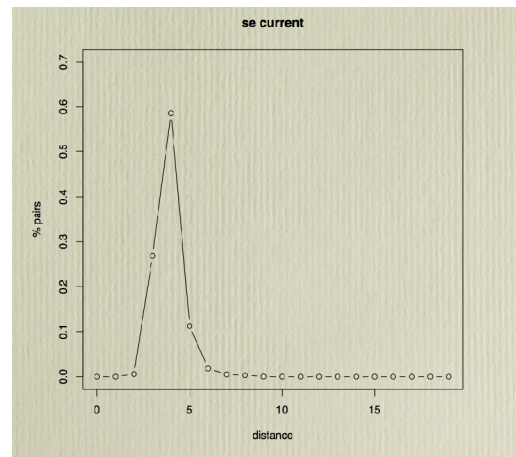
- **it** : only italian users
- **se** : only swedish users
- **it + se** : only italian and swedish users
- **us** : only US users
- the whole facebook (750m nodes)

based on users current geo-IP location

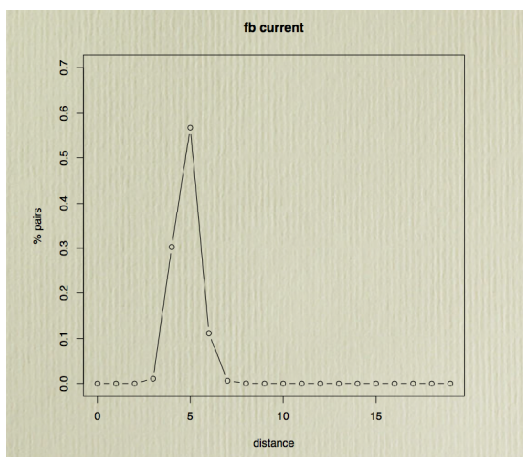
distance distribution (it)



distance distribution (se)



distance distribution (fb)



average distance

	2008	2012
it	6.58	3.90
se	4.33	3.89
it+se	4.90	4.16
us	4.74	4.32
fb	5.28	4.74

fb 2012 : 92% pairs are reachable!

effective diameter

	2008	2012
it	9.0	5.2
se	5.9	5.3
it+se	6.8	5.8
us	6.5	5.8
fb	7.0	6.2

actual diameter

	2008	2012
it	> 29	= 25
se	> 16	= 25
it+se	> 21	= 27
us	> 17	= 30
fb	> 17	> 58

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Separating You and Me? 4.74 Degrees

By JOHN MARKOFF and SOMINI SENGUPTA
Published: November 21, 2011

The world is even smaller than you thought.

Enlarge This Image

Adding a new chapter to the research that cemented the phrase "six degrees of separation" into the language, scientists at Facebook and the University of Milan reported on Monday that the average number of acquaintances separating any two people in the world was not six but 4.74.

The original "six degrees" finding, published in 1967 by the psychologist Stanley Milgram, was drawn from 296 volunteers who were asked to send a message by postcard, through friends and then friends of friends, to a specific person in a Boston suburb.

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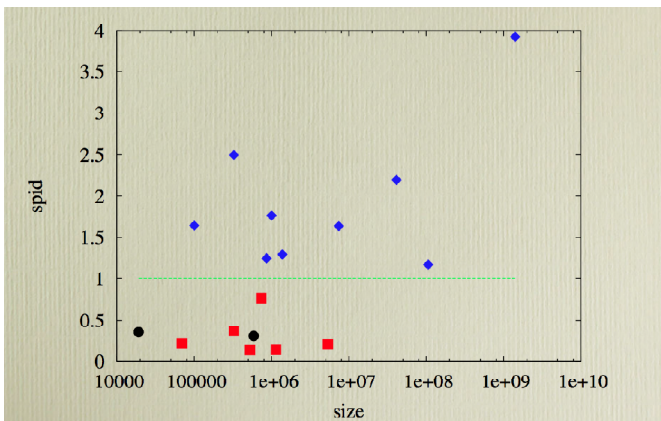
Billioe håndværkermobil
Flere fordelagtige abonnemønter. Bestil din håndværkermobil

another application : spid

[Boldi et al., 2011]

- **spid** : shortest-paths index of dispersion
- the **ratio** between **variance** and **average** in the distance distribution
- **spid** < 1 : the distribution is subdispersed
- **spid** > 1 : is superdispersed
- **web graphs** and **social networks** have **different** spid!

spid plot



the spid conjecture

- [Boldi et al., 2011] **conjecture** that spid is able to tell social networks from web graphs
- **average distance** alone would **not** changeable and depends on the scale
- spid, instead, seems to have a **clear cutpoint** at 1
- what is facebook spid?

the spid conjecture

- [Boldi et al., 2011] **conjecture** that spid is able to tell social networks from web graphs
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0.093

indexing distances in large graphs

shortest-path distances in large graphs

- **input:** consider a graph $G = (V, E)$
- and nodes s and t in V
- **goal:** compute the shortest-path distance $d(s, t)$ from s to t
- do it **very fast**

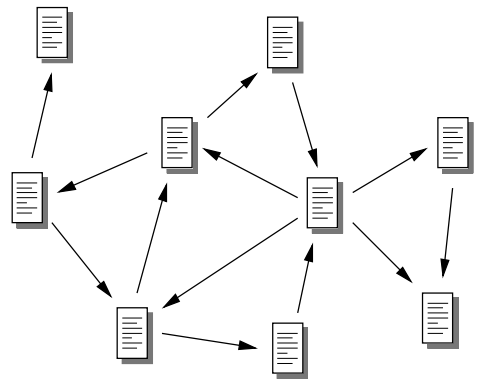
well-studied problem

different strategies

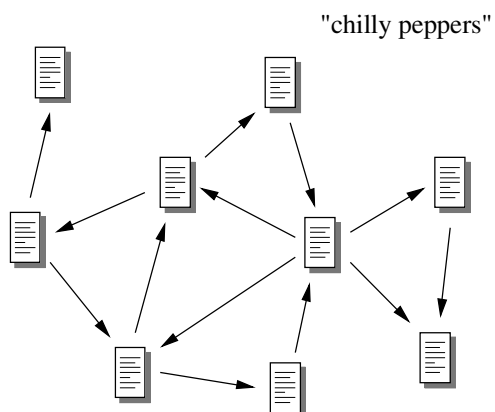
- **lazy**
 - compute shortest path at **query time**
 - Dijkstra, BFS
 - no precomputation
 - BFS takes $O(m)$
 - **too expensive** for large graphs
- **eager**
 - **precompute** all-pairs shortest paths
 - Floyd-Warshall, matrix multiplication
 - $O(n^3)$ precomputation, $O(n^2)$ storage
 - **too large** to store

applications of shortest-path queries

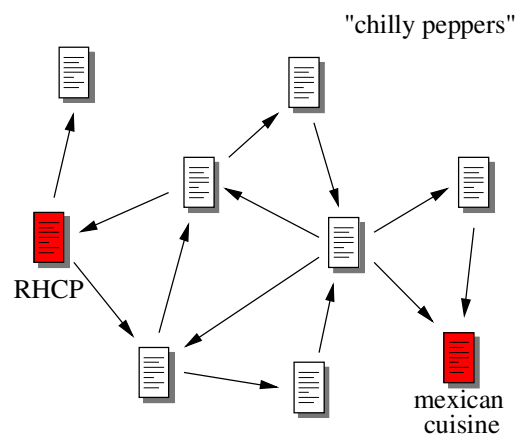
searching in graphs — I. context-sensitive search



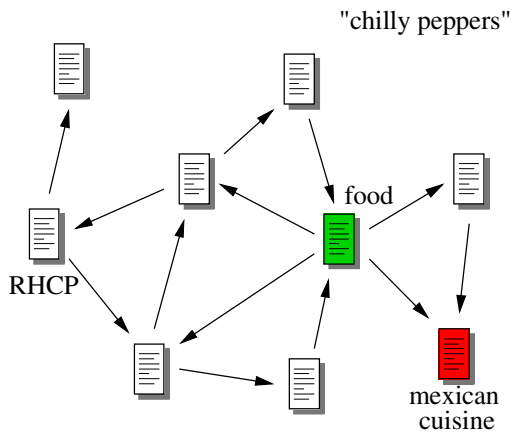
searching in graphs — I. context-sensitive search



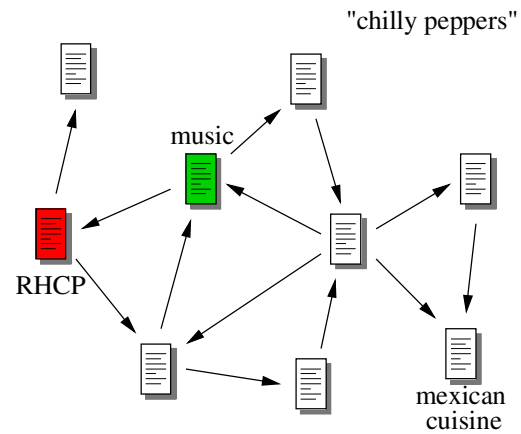
searching in graphs — I. context-sensitive search



searching in graphs — I. context-sensitive search



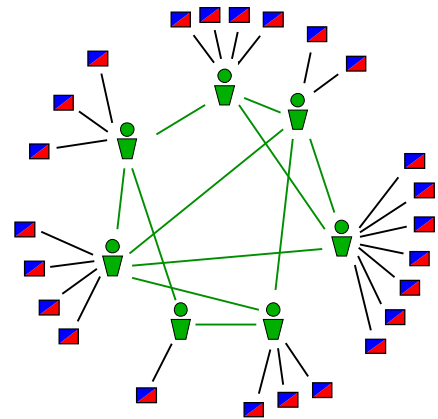
searching in graphs — I. context-sensitive search



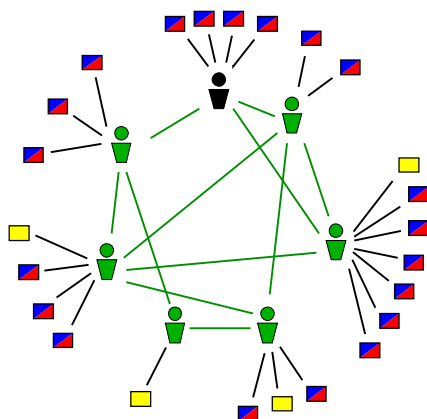
searching in graphs — I. context-sensitive search

- customize search results to the user's current page or recent history of pages have visited
- increasing relevance of answers
- disambiguation
- suggesting links to wikipedia editors

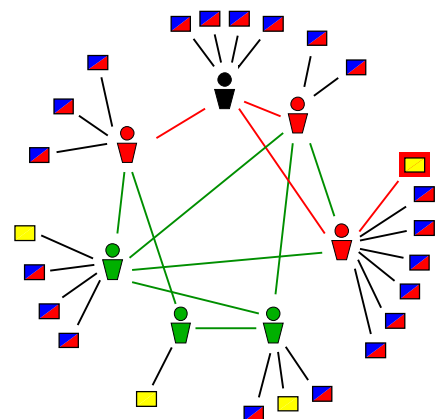
searching in graphs — II. social search



searching in graphs — II. social search



searching in graphs — II. social search



searching in graphs — II. social search

- consider more information than just contacts
 - preferences
 - geographical information
 - comments
 - favorites
 - tags
 - etc.

machine-learning approach

- learn a **ranking function** that **combines** a **large** number of **features**
 - content-based features:**
 - TF/IDF, BM25, etc., as in traditional IR and web search
 - content similarity between the querying node and a target node
 - link-based features:**
 - PageRank
 - **shortest-path distance** from the querying node to a target node
 - spectral distance from the querying node to a target node
 - graph-based similarity measures
 -

well-studied problem

different strategies

- **lazy**
 - compute shortest path at **query time**
 - Dijkstra, BFS
 - no precomputation
 - BFS takes $O(m)$
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 - **too large** to store

anything in between?

- is there a **smooth tradeoff** between $\langle O(1), O(m) \rangle$ and $\langle O(n^2), O(1) \rangle$

distance oracles

[Thorup and Zwick, 2005]

- given a graph $G = (V, E)$
- an (α, β) -**approximate distance oracle** is a **data structure** S that
- for a query pair of nodes (u, v) , S returns $d_S(u, v)$ s.t.

$$d(u, v) \leq d_S(u, v) \leq \alpha d(u, v) + \beta$$

- α called **stretch** or **distortion**
- consider the **preprocessing time**, the **required space**, and the **query time**

distance oracles

[Thorup and Zwick, 2005]

- given k , construct an oracle with storage $O(kn^{1+1/k})$, query time $O(k)$, stretch $2k - 1$
- $k = 1$
 \Rightarrow APSP
- $k = \log n$
 \Rightarrow storage $O(n \log n)$, query time $O(\log n)$, stretch $O(\log n)$

distance oracles — preprocessing

[Das Sarma et al., 2010]

- 1 $r = \lfloor \log |V| \rfloor$
- 2 sample $r + 1$ sets of sizes $1, 2, 2^2, 2^3, \dots, 2^r$
- 3 call the sampled sets S_0, S_1, \dots, S_r
- 4 for each node u and each set S_i compute (w_i, δ_i) ,
where $\delta_i = d(u, w_i) = \min_{v \in S_i} \{d(u, v)\}$
- 5 $\text{SKETCH}[u] = \{(w_0, \delta_0), \dots, (w_r, \delta_r)\}$
- 6 repeat k times

distance oracles — query processing

[Das Sarma et al., 2010]

given query (u, v)

- 1 obtain $\text{SKETCH}[u]$ and $\text{SKETCH}[v]$
- 2 w in $\text{SKETCH}[u]$ and $\text{SKETCH}[v]$
- 3 for each common node w , compute $d(u, w)$ and $d(w, v)$
- 4 return the minimum of $d(u, w) + d(w, v)$,
taken over all common node w 's
- 5 if no common w is present, then return ∞

landmark-based approach

- precompute: $d(s, l)$ for all $s, l \in V$
- then
 $|d(s, l) - d(t, l)| \leq d(s, t) \leq d(s, l) + d(l, t)$
- precompute: distances to d landmarks, l_1, \dots, l_d
 $\max_i |d(s, l_i) - d(t, l_i)| \leq d(s, t) \leq \min_i (d(s, l_i) + d(l_i, t))$
- obtain a range estimate in time $O(d)$ (i.e., constant)

landmark-based approach

- motivated by indexing general metric spaces
- used for estimating latency in the internet
[Ng and Zhang, 2008]
- typically randomly chosen landmarks

theoretical results

[Kleinberg et al., 2004]

- random landmarks can provide distance estimates with distortion $(1 + \delta)$ for a fraction of at least $(1 - \epsilon)$ of pairs
- number of landmarks required depends on ϵ , δ , and the doubling dimension k of the metric space

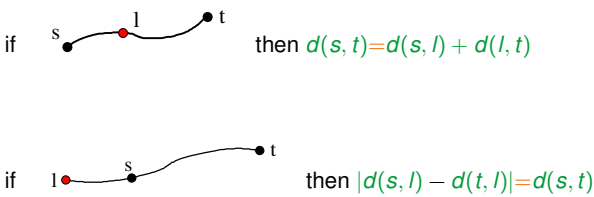
approximation guarantee in practice

what does a logarithmic approximation guarantee mean in a small-world graph?

the landmark selection problem

how to choose good landmarks in practice?

good landmarks



good (upper-bound) landmarks

- a landmark l covers a pair (s, t) if l is on a shortest path from s to t
- problem definition: $L \subseteq V$ of k landmarks that cover as many pairs $(s, t) \in V \times V$ as possible
- NP-hard
- for $k = 1$: the node with the highest centrality betweenness
- for $k > 1$: apply a “natural” set-cover approach (but $O(n^3)$)

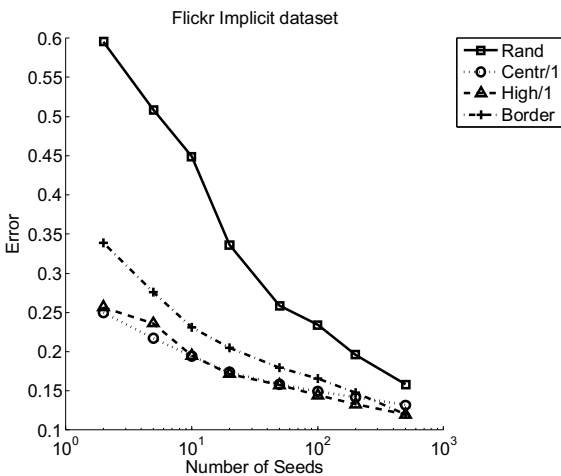
landmark selection heuristics

- high-degree nodes
- high-centrality nodes
- “constrained” versions
 - once a node is selected none of its neighbors is selected
- “clustered” versions
 - cluster the graph and select one landmark per cluster
 - select landmarks on the “borders” between clusters

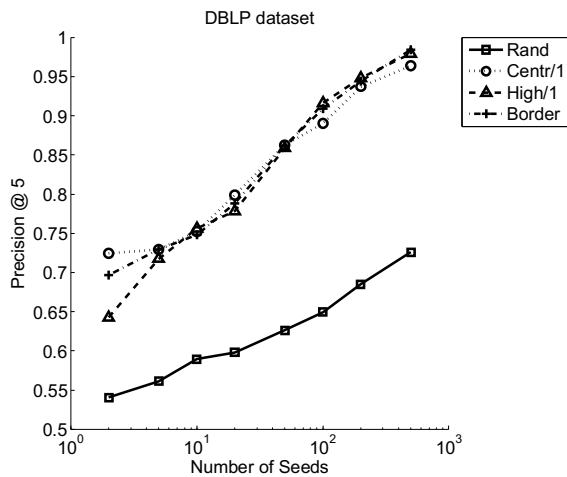
datasets

	# nodes	# edges	median distance	effective diameter	clustering coefficient
DBLP	226 K	716 K	9	13	0.47

flickr-implicit — distance error

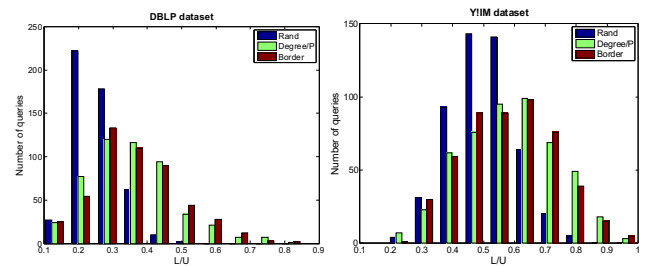


DBLP — precision @ 5



triangulation task

[Kleinberg et al., 2004]



comparing with exact algorithm

[Goldberg and Harrelson, 2005]

landmarks (10%)	Fl.-E	Fl.-I	Wiki	DBLP	YIM
Method	CENT	CENT	CENT/P	BORD/P	BORD/P
Landmarks used	20	100	500	50	50
Nodes visited	1	1	1	1	1
Operations	20	100	500	50	50
CPU ticks	2	10	50	5	5
ALT (exact)	Fl.-E	Fl.-I	Wiki	DBLP	YIM
Method	Ikeda	Ikeda	Ikeda	Ikeda	Ikeda
Landmarks used	8	4	4	8	4
Nodes visited	7245	10337	19616	2458	2162
Operations	56502	41349	78647	19666	8648
CPU ticks	7062	10519	25868	1536	1856

acknowledgements



Paolo Boldi



Charalampos Tsourakakis






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