Locality-sensitive hashing and its applications

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ACM Kanellakis Award 2013

A frequent issue

Given U users, described with a set of **d** features, the goal is to find (the largest) group of *similar* users

Features = Personal data, preferences, purchases,
 navigational behavior, search behavior, followers/ing,

Similareitty (en is, type) casty and unantionath altyetakiemy the reset of features of users u1 and u2, returns a value in [0,1]

Users could also be Web pages (depp), products (recommendation tweets/news/search results (visualization)

000110010

100110010

Solution #1

Try all *groups of users* and, for each group, check the (average) similarity among all its users.

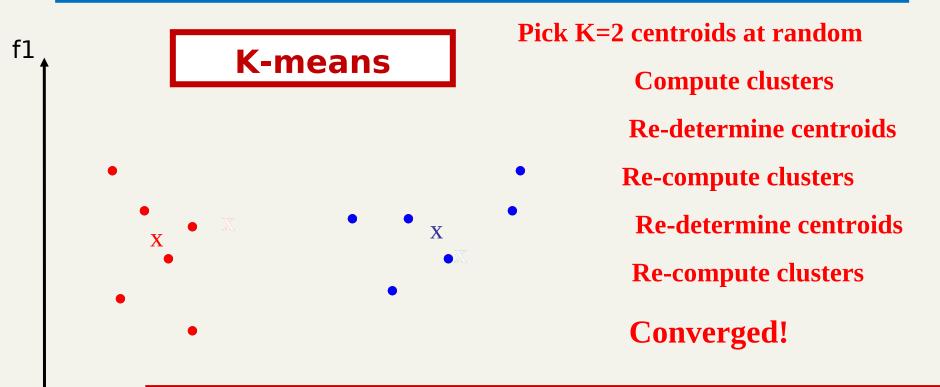
```
# Sim computations \cong 2^{U} \times U^{2}
In the case of Facebook this is > 2^{1billion} \times (10^{9})^{2}
If we limit groups to have a size \leq L users

# Sim computations \cong U^{L} \times L^{2}
(Even if 1ns/sim and L=10, it is > (10^{9})^{10}/10^{9} secs > 10^{70} \times 2^{15})
```

No faster CPU/GPU, multi-cores,... could help!

Solution #2: introduce approximation

Interpret every user as a point in a **d**-dim space, and then apply a *clustering* algorithm



Each iteration takes $\cong K \times U$

Solution #2: few considerations

- Cost per iteration = $\mathbf{K} \times \mathbf{U}$, #iterations is typically small
- What about optimality? It is locally optimal [recently, some researchers showed how to introduce some guarantee]
- What about the Sim-cost? Comparing users/points costs Θ(d) in time and space [notice that d may be bi/mile Text
 - In T time, we can manage $U = T^{1/3}$
- [≅y] Using s-faster CPU ≈ using sT time an old CPU

Solution #3: introduce randomization

Generate a **fingerprint** for every user that is **much shorter** than **d** and allows to transform similarity into **equality** of fingerprints.

- ✓ It is randomized, correct with high probability
- It guarantees *local access* to data, which is good for speed in disk/distributed setting

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A warm-up problem

- Consider vectors p,q of d binary features
- Hamming distance D(p,q) = #bits where p and q differ
- Define hash function h by choosing a set l of k random coordinates

h(p) = projection of vector p on l's coordinates

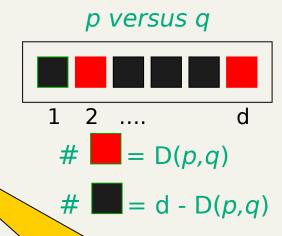
Example: Pick $I = \{1,4\}$ (k=2), then h(p=01011) = 01

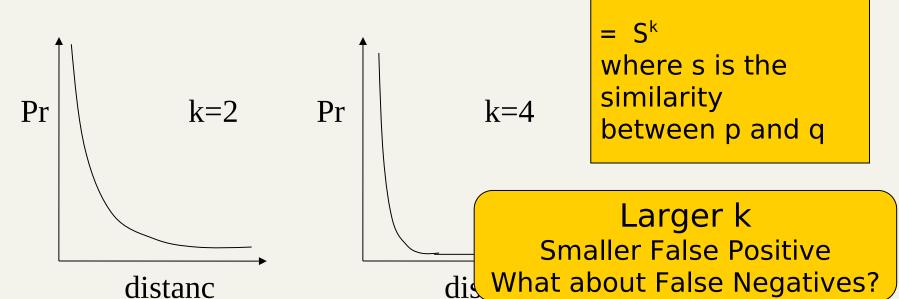
A key property

Pr[picking x s.t. p[x]=q[x]]= (d - D(p,q))/d

$$\Pr[h(p) = h(q)] = (1 - \frac{D(p,q)}{d})^k$$

We can vary the probability by changing





Larger L Smaller False Negatives

Reiterate L times

- 1) Repeat **L times** the **k-projections** h_i(p)
- 2) We set $g(p) = \langle h_1(p), h_2(p), ..., h_l(p) \rangle \frac{\text{Sketch}(p)}{\text{Sketch}(p)}$
- 3) Declare «p matches q» if **at least** one $h_i(p)=h_i(q)$

Example:

```
We set k=2, L=3, let p=01001 and q=01101
```

•
$$11 = \{3,4\}$$
, we have $h_1(p) = 00$ and $h_1(q)=10$

•12 = {1,3}, we have
$$h_2(p) = 00$$
 and $h_2(q)=01$

•13 = {1,5}, we have
$$h_3(p) = 01$$
 and $h_3(q) = 01$ p and q declared

p and q declared to match!!

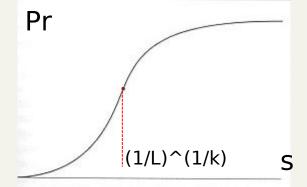
Measuring the error probability

$$\Pr[h_i(p) = h_i(q)] = (1 - \frac{D(p,q)}{d})^k = s^k$$

The g() consists of L independent hashes hi

$$Pr[g(p) \text{ matches } g(q)] = 1 - Pr[h_i(p) \neq h_i(q), \forall i=1, ...,$$

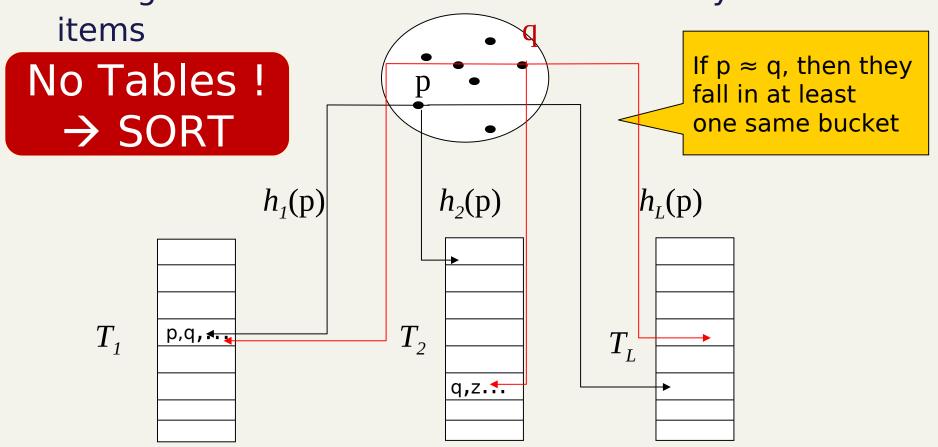
$$\Pr(g(p) \cong g(q)) = 1 - (1 - s^k)^L$$



The case: Groups of similar items

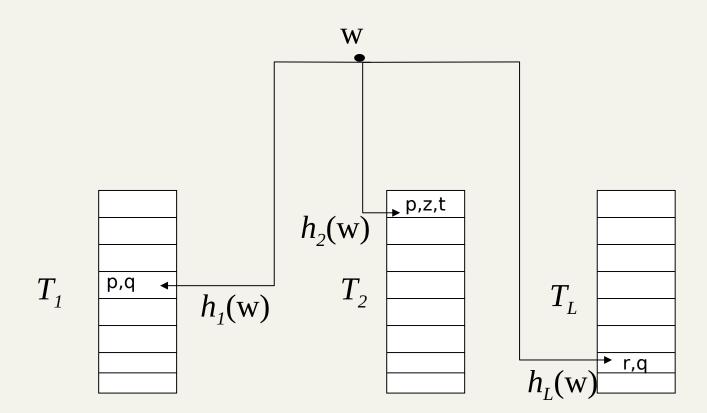
Buckets provide the candidate similar items

«Merge» similar sets over L rounds if they share



The case of on-line query

Given a query w, find the similar indexed vectors: check the vectors in the buckets $h_i(w)$ for all j=1,...,L



LSH versus K-means

- What about optimality? K-means is locally optimal [LSH finds correct clusters with high probability]
- What about the Sim-cost? K-means compares vectors of d components [LSH compares very short (sketch) vectors]
- What about the cost per iteration? Typically K-means requires few iterations, each costs K × U ×
 d [LSH sorts U short items, few scans]
- What about K? In principle You could apply K-means over LSH-sketch vectors !!