Data Structures and Algorithms Bloom Filters

CS 225 Brad Solomon

April 19, 2024



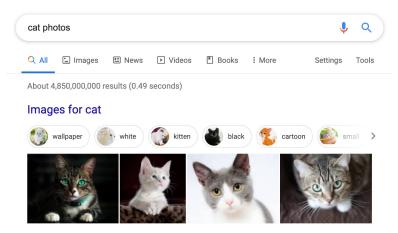
Department of Computer Science

Data Structures Review

What method would you use to build a search index on a collection of objects?

What method would you use to build a search index on a collection of objects *in a memory-constrained environment*?

Constrained by Big Data (Large N)

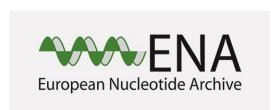


Google Index Estimate: >60 billion webpages

Google Universe Estimate (2013): >130 trillion webpages

What method would you use to build a search index on a collection of objects *in a memory-constrained environment*?

Constrained by Big Data (Large N)









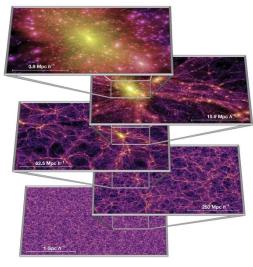
SRA

Sequence Read Archive (SRA) makes biological sequence data available to the research community to enhance reproducibility and allow for new discoveries by comparing data sets. The SRA stores raw sequencing data and alignment information from high-throughput sequencing platforms, including Roche 454 GS System®, Illumina Genome Analyzer®, Applied Biosystems SOLiD System®, Helicos Heliscope®, Complete Genomics®, and Pacific Biosciences SMRT®.

Sequence Read Archive Size: >60 petabases (1015)

What method would you use to build a search index on a collection of objects *in a memory-constrained environment*?

Constrained by Big Data (Large N)



| Sky Survey Projects | Data Volume |
|--|-------------------|
| DPOSS (The Palomar Digital Sky Survey) | 3 TB |
| 2MASS (The Two Micron All-Sky Survey) | 10 TB |
| GBT (Green Bank Telescope) | 20 PB |
| GALEX (The Galaxy Evolution Explorer) | 30 TB |
| SDSS (The Sloan Digital Sky Survey) | 40 TB |
| SkyMapper Southern Sky Survey | 500 TB |
| PanSTARRS (The Panoramic Survey Telescope and Rapid Response System) | ~ 40 PB expected |
| LSST (The Large Synoptic Survey Telescope) | ~ 200 PB expected |
| SKA (The Square Kilometer Array) | ~ 4.6 EB expected |
| | |

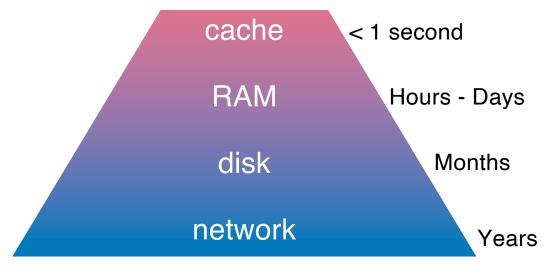
Table: http://doi.org/10.5334/dsj-2015-011

Estimated total volume of one array: 4.6 EB

Image: https://doi.org/10.1038/nature03597

What method would you use to build a search index on a collection of objects *in a memory-constrained environment*?

Constrained by resource limitations



(Estimates are Time x 1 billion courtesy of https://gist.github.com/hellerbarde/2843375)

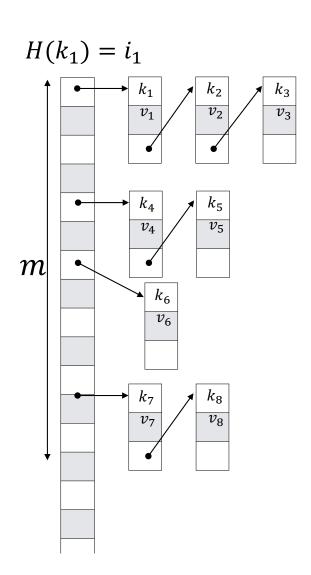
What method would you use to build a search index on a collection of objects *in a memory-constrained environment*?

Reducing storage costs

1) Throw out information that isn't needed

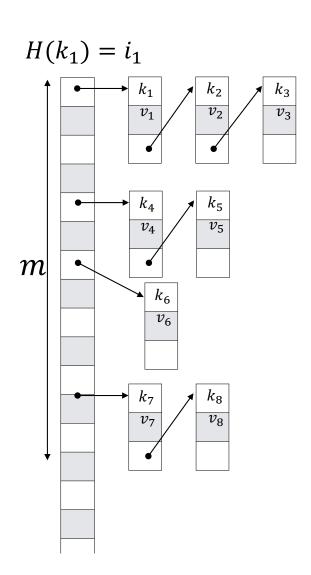
2) Compress the dataset

What can we remove from a hash table?



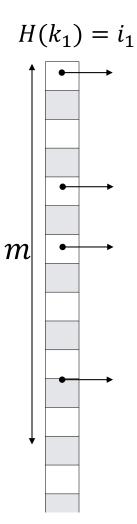
What can we remove from a hash table?

Take away values



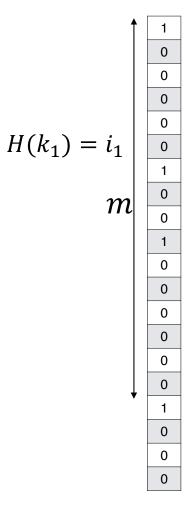
What can we remove from a hash table?

Take away values and keys



What can we remove from a hash table?

Take away values and keys



Bloom Filter: Insertion

 $S = \{ 16, 8, 4, 13, 29, 11, 22 \}$

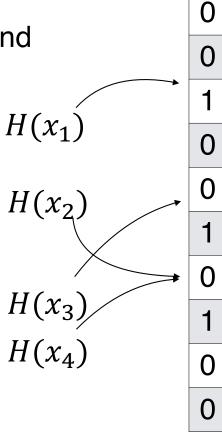
h(k) = k % 7

- 0 0
- 1 0
- 2 0
- 3 0
- 4 0
- 5 0
- 6 0

Bloom Filter: Insertion

An item is inserted into a bloom filter by hashing and then setting the hash-valued bit to 1

If the bit was already one, it stays 1



Bloom Filter: Deletion

```
S = { 16, 8, 4, 13, 29, 11, 22 } __delete(13)
```

h(k) = k % 7

 $0 \quad 0$

1 1

2 1

3 0

4 1

5 0

6 1

delete(29)

Bloom Filter: Search

```
S = \{ 16, 8, 4, 13, 29, 11, 22 \} _find(16)
h(k) = k % 7
```

_find(20)

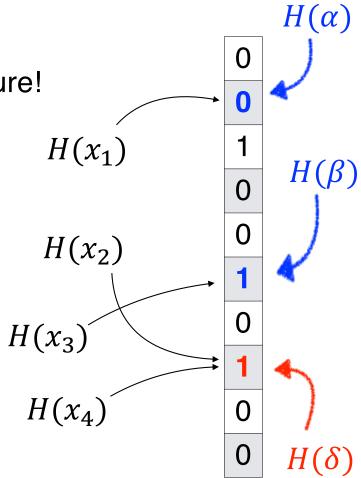
_find(3)

Bloom Filter: Search

The bloom filter is a *probabilistic* data structure!

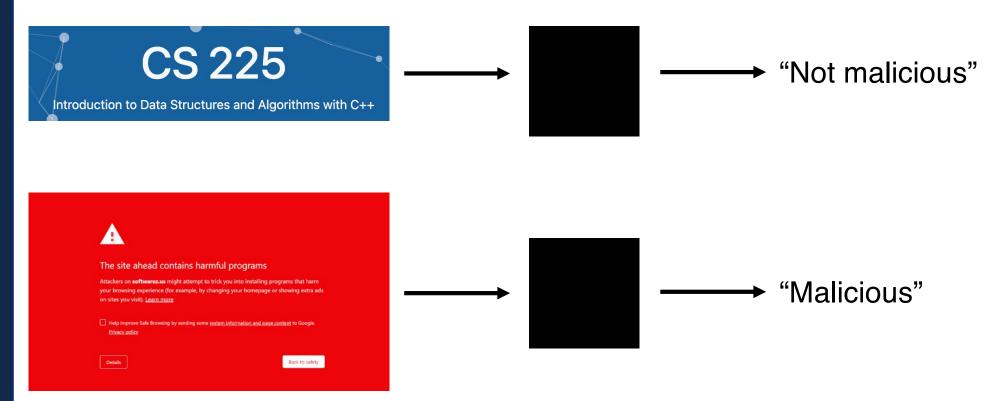
If the value in the BF is 0:

If the value in the BF is 1:



Probabilistic Accuracy: Malicious Websites

Imagine we have a detection oracle that identifies if a site is malicious



Probabilistic Accuracy: Malicious Websites

Imagine we have a detection oracle that identifies if a site is malicious

True Positive:

False Positive:

False Negative:

True Negative:

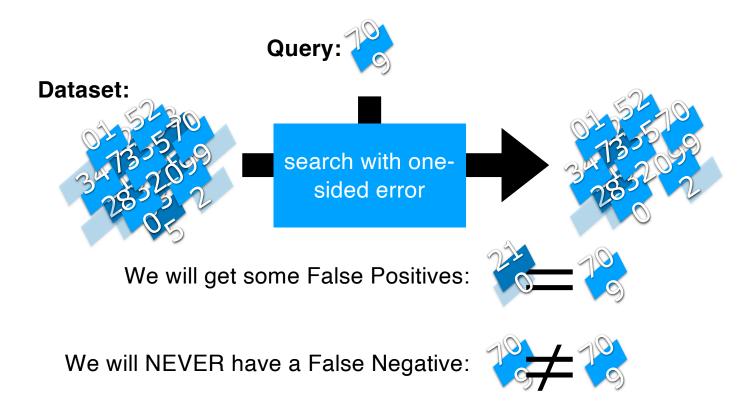
Imagine we have a **bloom filter** that **stores malicious sites...**

Bit Value = 1 Bit Value = 0 H(z)H(z)'Yes' 'No' True Positive False Negative H(z)H(z)'Yes' 'No' True Negative False Positive

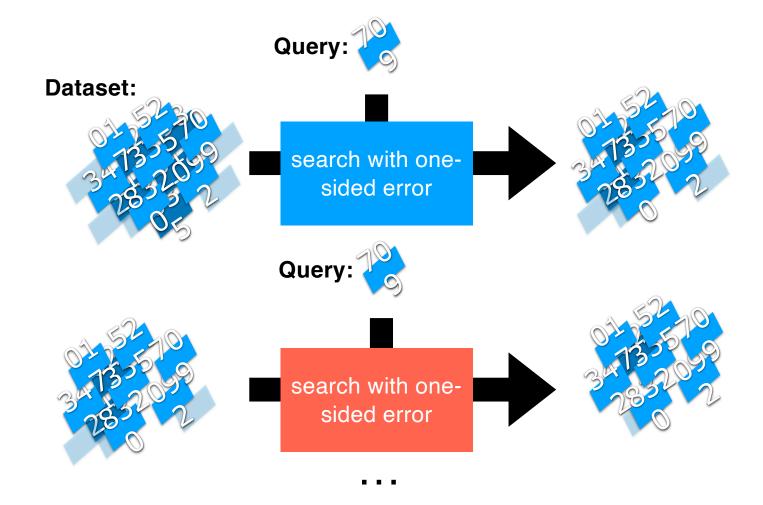
Item Inserted

Item NOT inserted

Probabilistic Accuracy: One-sided error

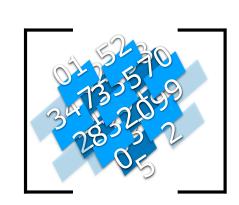


Probabilistic Accuracy: One-sided error



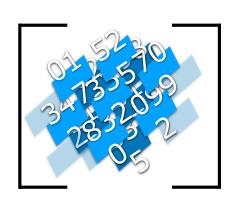
Use many hashes/filters; add each item to each filter

 h_1



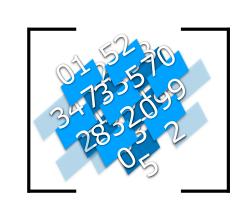
Use many hashes/filters; add each item to each filter

 h_2



Use many hashes/filters; add each item to each filter

| | 0 | | 0 | | 0 |
|---------------|---|--------------------|---|-------|---|
| | 1 | | 0 | | 1 |
| | 0 | | 0 | | 1 |
| | 1 | | 1 | | 1 |
| | 0 | | 0 | | 0 |
| | 0 | | 0 | | 0 |
| | 0 | | 0 | | 1 |
| | 1 | | 1 | | 1 |
| h_1 | 0 | h_2 | 0 | 1_ | 0 |
| $'^{\iota_1}$ | 1 | $^{\prime\prime}2$ | 1 | h_3 | 1 |
| | 1 | | 0 | | 1 |
| | 0 | | 0 | | 0 |
| | 1 | | 1 | | 1 |
| | 0 | | 1 | | 0 |
| | 1 | | 1 | | 1 |
| | 1 | | 0 | | 1 |
| | 0 | | 0 | | 0 |
| | 1 | | 1 | | 1 |
| | 0 | | 0 | | 0 |
| | 1 | | 0 | | 1 |
| | | 1 | | | |



Use many hashes/filters; add each item to each filter

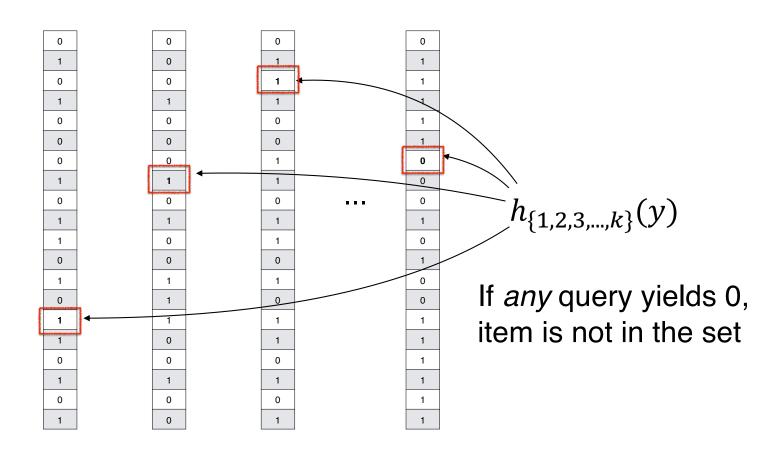
| | 0 | | 0 | | 0 | | 0 |
|----------------|---|-------|---|-------|---|-----------|---|
| | 1 | | 0 | | 1 | | 1 |
| | 0 | | 0 | | 1 | | 1 |
| | 1 | | 1 | | 1 | | 1 |
| | 0 | | 0 | | 0 | | 1 |
| | 0 | | 0 | | 0 | | 1 |
| | 0 | | 0 | | 1 | | 0 |
| | 1 | | 1 | | 1 | | 0 |
| h_1 | 0 | h_2 | 0 | h_3 | 0 | h_k | 0 |
| $\iota\iota_1$ | 1 | n_2 | 1 | 113 | 1 | r_k | 1 |
| | 1 | | 0 | | 1 | | 0 |
| | 0 | | 0 | | 0 | | 1 |
| | 1 | | 1 | | 1 | | 0 |
| | 0 | | 1 | | 0 | | 0 |
| | 1 | | 1 | | 1 | | 1 |
| | 1 | | 0 | | 1 | | 1 |
| | 0 | | 0 | | 0 | | 1 |
| | 1 | | 1 | | 1 | | 1 |
| | 0 | | 0 | | 0 | | 1 |
| | 1 | | 0 | | 1 | | 1 |

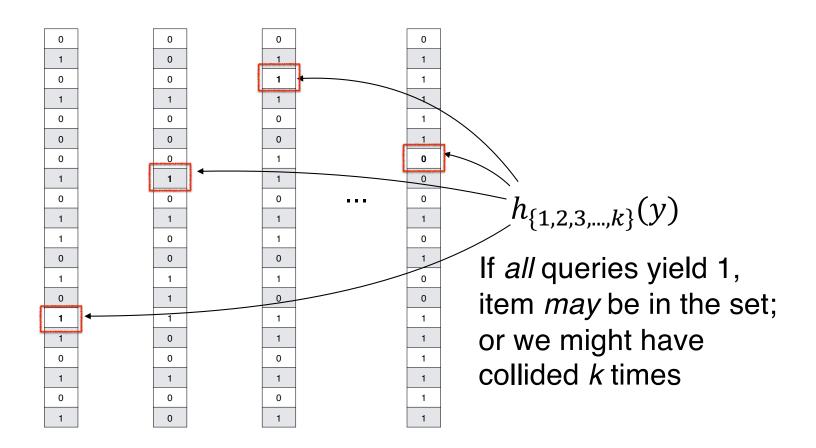
| 0 | |
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| 0 | |
| 1 | |

$$h_{\{1,2,3,...,k\}}(y)$$





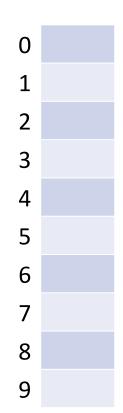
Using repeated trials, even a very bad filter can still have a very low FPR!

If we have k bloom filter, each with a FPR p, what is the likelihood that **all** filters return the value '1' for an item we didn't insert?

But doesn't this hurt our storage costs by storing k separate filters?

| | 0 | | 0 | | 0 | | | 0 |
|-------|---|-------|---|-------|---|---|-------|---|
| | 1 | | 0 | | 1 | | | 1 |
| | 0 | | 0 | | 1 | | | 1 |
| | 1 | | 1 | | 1 | | | 1 |
| | 0 | | 0 | | 0 | | | 1 |
| | 0 | | 0 | | 0 | | | 1 |
| | 0 | | 0 | | 1 | | | 0 |
| | 1 | | 1 | | 1 | | | 0 |
| h | 0 | h_2 | 0 | h_3 | 0 | | h _ | 0 |
| h_1 | 1 | 1 12 | 1 | 113 | 1 | | h_k | 1 |
| | 1 | | 0 | | 1 | | | 0 |
| | 0 | | 0 | | 0 | | | 1 |
| | 1 | | 1 | | 1 | | | 0 |
| | 0 | | 1 | | 0 | | | 0 |
| | 1 | | 1 | | 1 | | | 1 |
| | 1 | | 0 | | 1 | | | 1 |
| | 0 | | 0 | | 0 | | | 1 |
| | 1 | | 1 | | 1 | | | 1 |
| | 0 | | 0 | | 0 | | | 1 |
| | 1 | | 0 | | 1 | | | 1 |
| | | , | | | | • | ' | |

Rather than use a new filter for each hash, one filter can use k hashes



$$S = \{ 6, 8, 4 \}$$

$$h_1(x) = x \% 10$$
 $h_2(x) = 2x \% 10$ $h_3(x) = (5+3x) \% 10$

$$h_3(x) = (5+3x) \% 10$$

9

Rather than use a new filter for each hash, one filter can use k hashes

```
0 0 h<sub>1</sub>(x) = x % 10 h<sub>2</sub>(x) = 2x % 10 h<sub>3</sub>(x) = (5+3x) % 10

1 0
2 1
3 1
4 1
5 0
6 1
7 1
8 1
```

Bloom Filter

A probabilistic data structure storing a set of values

 $H = \{h_1, h_2, \dots, h_k\}$

- 0
- 0
- 1
- 0
- 0
- 1
- 0
- _
- \cap
- 0

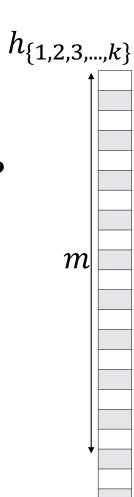
Built from a bit vector of length m and k hash functions

Insert / Find runs in: _____

Delete is not possible (yet)!

Given bit vector of size m and k SUHA hash function

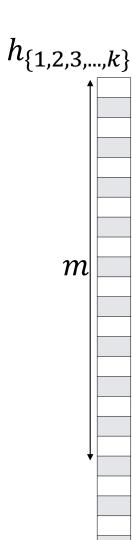
What is our expected FPR after n objects are inserted?



Given bit vector of size m and 1 SUHA hash function

What's the probability a specific bucket is 1 after one object is inserted?

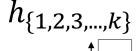
Same probability given k SUHA hash function?



Given bit vector of size m and k SUHA hash function

Probability a specific bucket is 0 after one object is inserted?

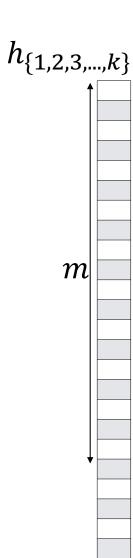
After n objects are inserted?



m

Given bit vector of size m and k SUHA hash function

What's the probability a specific bucket is ${\bf 1}$ after n objects are inserted?



 $h_{\{1,2,3,...,k\}}$

Given bit vector of size m and k SUHA hash function

What is our expected FPR after n objects are inserted?

The probability my bit is 1 after n objects inserted

$$\left(1-\left(1-\frac{1}{m}\right)^{nk}\right)^k$$

The number of [assumed independent] trials

m

Vector of size m, k SUHA hash function, and n objects

To minimize the FPR, do we prefer...

(A) large k

(B) small k

$$\left(1-\left(1-\frac{1}{m}\right)^{nk}\right)^k$$

