Data Structures and Algorithms Bloom Filters

CS 225 Brad Solomon November 18, 2024

Why store

when you (an store

K,V



Department of Computer Science

Announcements

MP_mosaic survey EC reached

MP_traversal survey EC **not reached** (Have until 11/20 to submit!)

MP_puzzle released, due after break. Break doesn't count as a week

Glab this wrek

Learning Objectives



Review when you would prefer different data structures

Build a conceptual understanding of a bloom filter

Review probabilistic data structures and one-sided error

Formalize the math behind the bloom filter $T_{widhesday}$

Which collision resolution strategy is better? Big Records: Open hushing (Seperate chaining) G (an pass by lef b closed hashing (ant allocate 94 this scale Structure Speed: (Josed hashing (Double hashing)

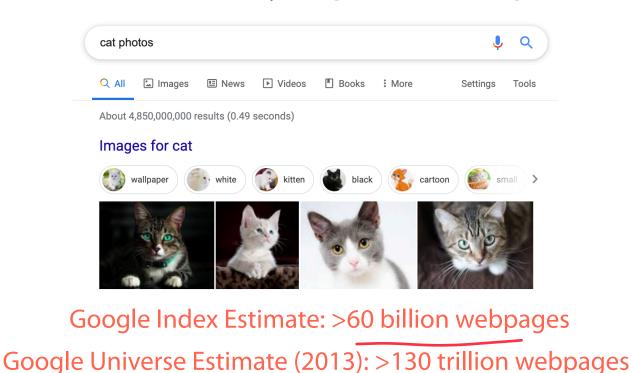
What structure do hash tables implement? Dictionaries

What constraint exists on hashing that doesn't exist with BSTs? () Probabilistic! QS imple unifour hashing assumption (SUHA) (3) Pseudo-amortized Why talk about BSTs at all? - SPL resize when Y < 1 (Some date is not hashable Sordered dates of useful (Nearest neighbor)

| Running Times | | y A | To Several To Several (KD Tree) | |
|---------------|--|------------------------|---------------------------------------|--|
| | Hash Table | AVL | Linked List | |
| Find | Expectation*: O(1)*** Worst Case: O(n) Expectation | O(log n) | O(n) | |
| Insert | Expectation*: O(1)*** Worst Case: O(n) | Guaranterd O(log n) | O(1) | |
| Storage Space | O(n) | O(n) | O(n) | |

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)



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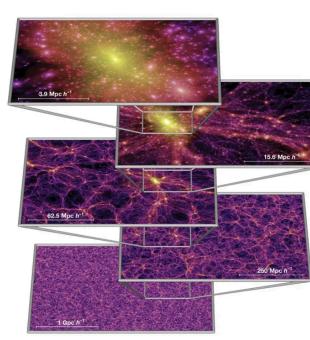
SRA

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)

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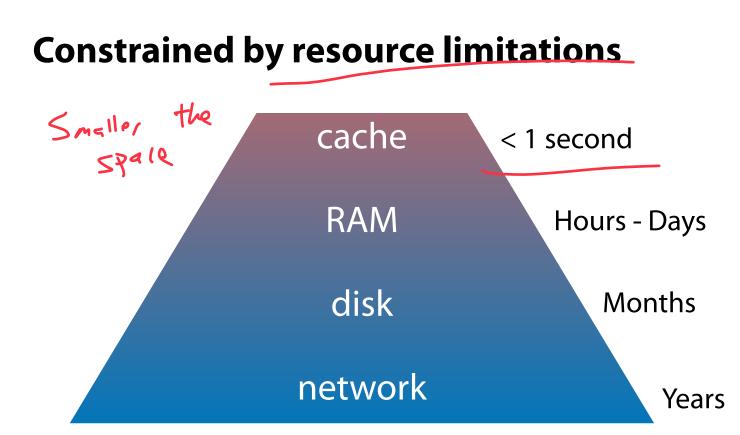
| Sky Survey Projects | Data Volume |
|--|-------------------|
| DPOSS (The Palomar Digital Sky Survey) | 3 TB |
| 2MASS (The Two Micron All-Sky Survey) | 10 TB |
| GBT (Green Bank Telescope) | 20 PB |
| GALEX (The Galaxy Evolution Explorer) | 30 TB |
| SDSS (The Sloan Digital Sky Survey) | 40 TB |
| 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*?



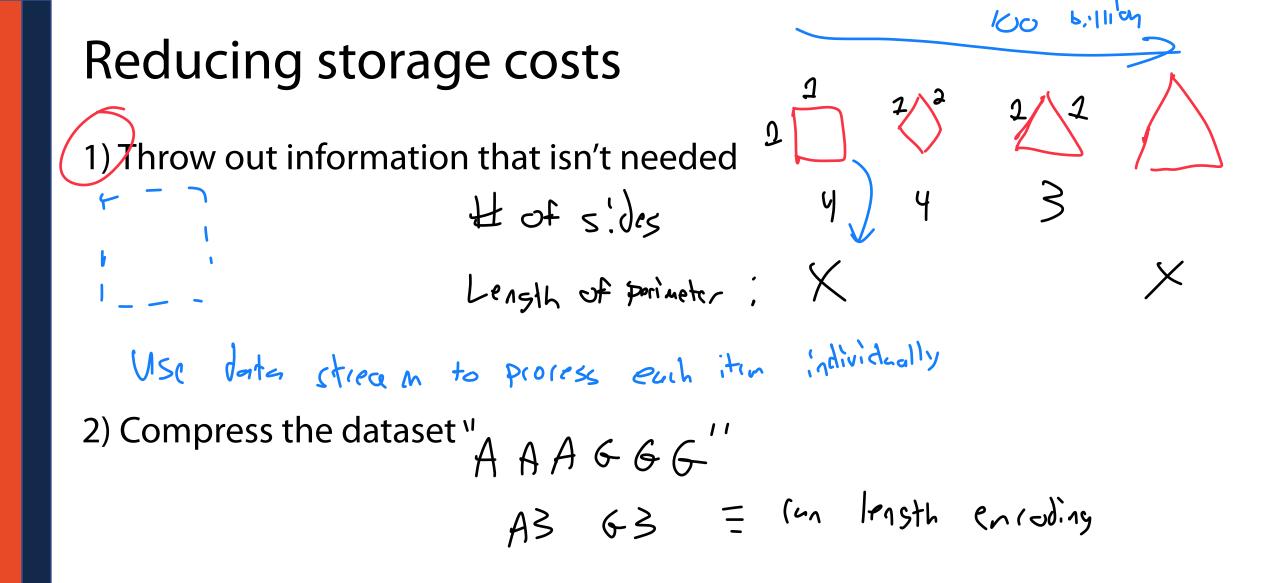
(Estimates are Time x 1 billion courtesy of https://gist.github.com/hellerbarde/2843375)

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

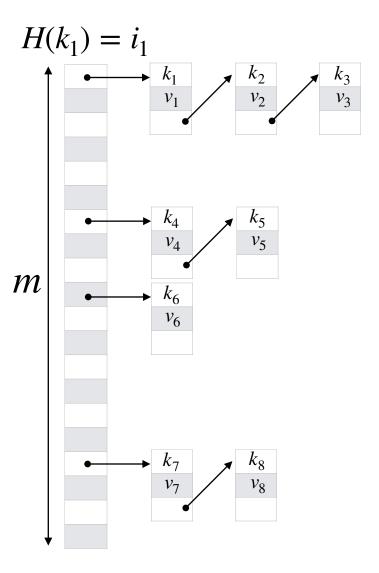
1) Make a space-efficient encoding (Compress the date)

3) Make a distributed network

Make - bloom filter

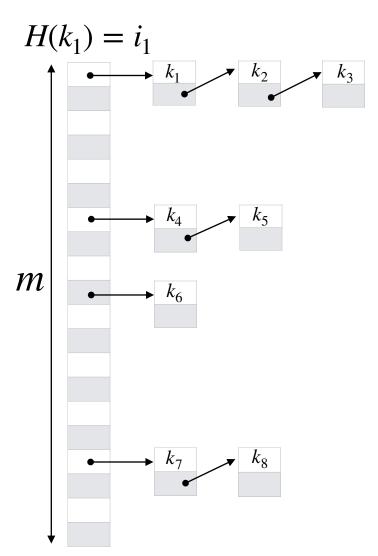


What can we remove from a hash table?



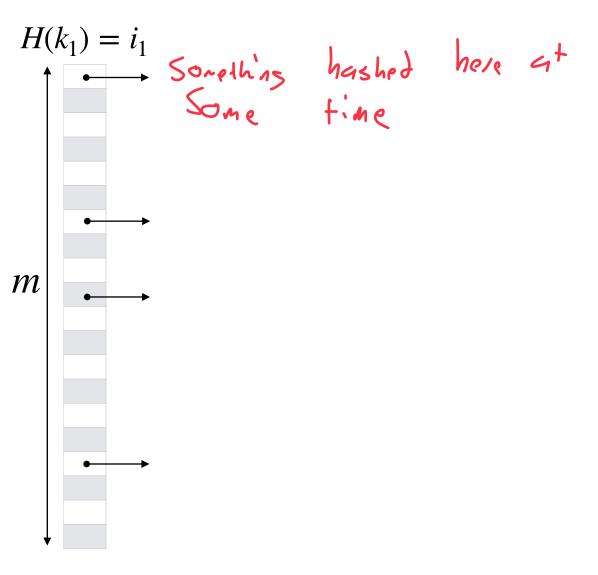
What can we remove from a hash table?

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

This is a **bloom filter**

 $H(k_1) = i_1$ 1 & Something hashed here 0 & Nothing hashed here 0 0 m 0 0 0 0 Storage (Small constant) Storage (1 bit per block)

Bloom Filter ADT

Constructor

Insert

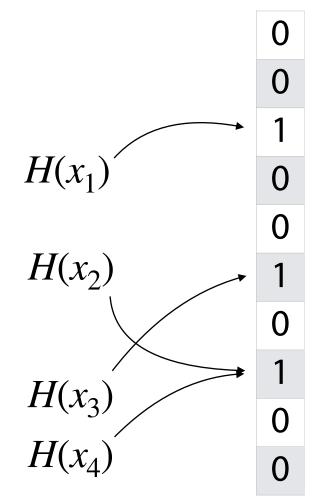
Find

Empty BF is vector lancy of Os **Bloom Filter: Insertion** 1) Hash Key to hash value (address) **S** = { 16, 8, 4, 13, 29, 11, 22 } h(k) = k % 72) Set bit to 1 at address 0 \leftarrow h(∂ 9)=1 ø1 1 No collision possible Ø1 2 3 0 I says "Something hashed of some point" Ø1 4 5 0 Ø 2 6

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 } h(k) = k % 70 1 0 2 1 3 0 1 4 5 0

Oh

10!

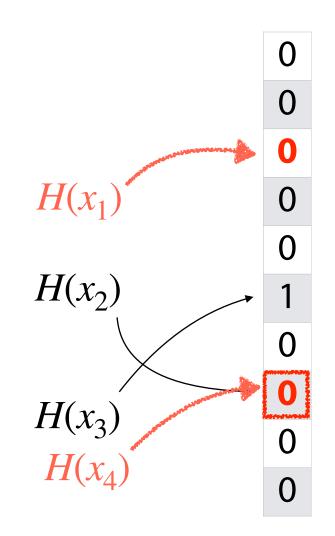
6

_delete(13) 1) Мисц Кеу 2) Syf b.7 40 O delete(29)

-find(8)

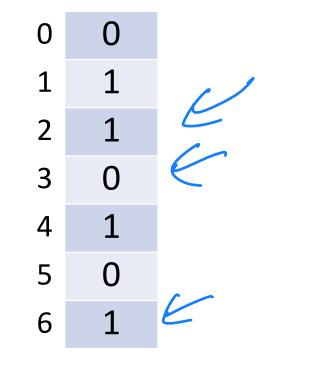
Bloom Filter: Deletion

Due to hash collisions and lack of information, items cannot be deleted!



Bloom Filter: Search

 $S = \{ 16, 8, 4, 13, 29, 11, 22 \}$ h(k) = k % 7 6



_find(16) > true! V 1) flash key 2) Look up value

_find(20) > The X 5 205 7= 6 False

_find(3) > False V (orrect

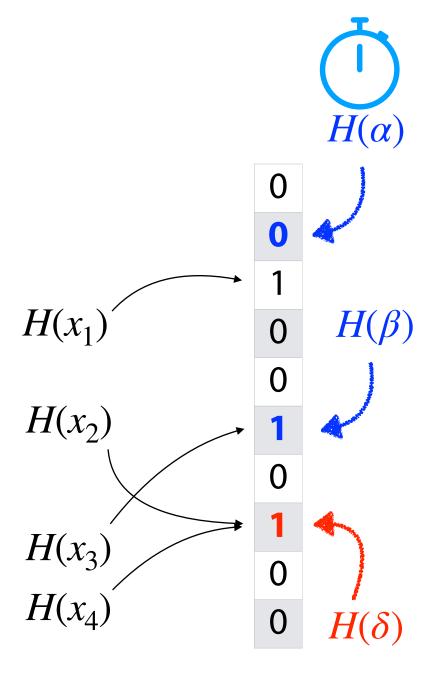
Probabilistic acluracy!

Bloom Filter: Search

The bloom filter is a *probabilistic* data structure!

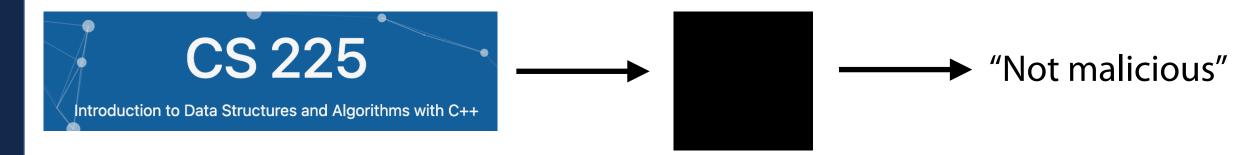
If the value in the BF is 0: 100% of the time item not in Set

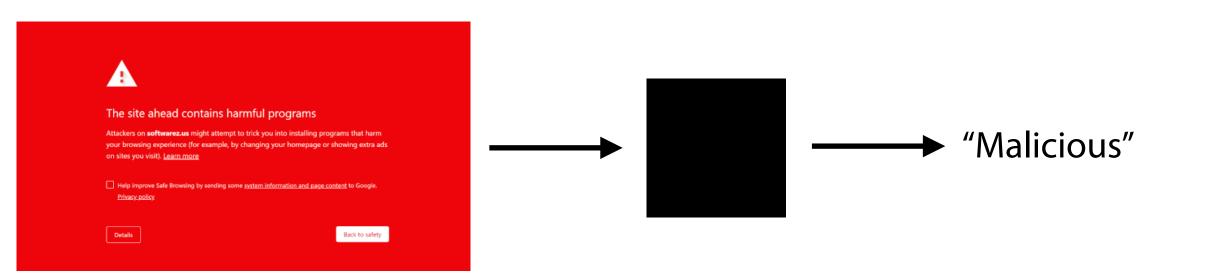
If the value in the BF is 1: item Might be in spt or is a hash collision



Probabilistic Accuracy: Malicious Websites

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





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True Positive: Oracle Says M

[actual webs; ite is m

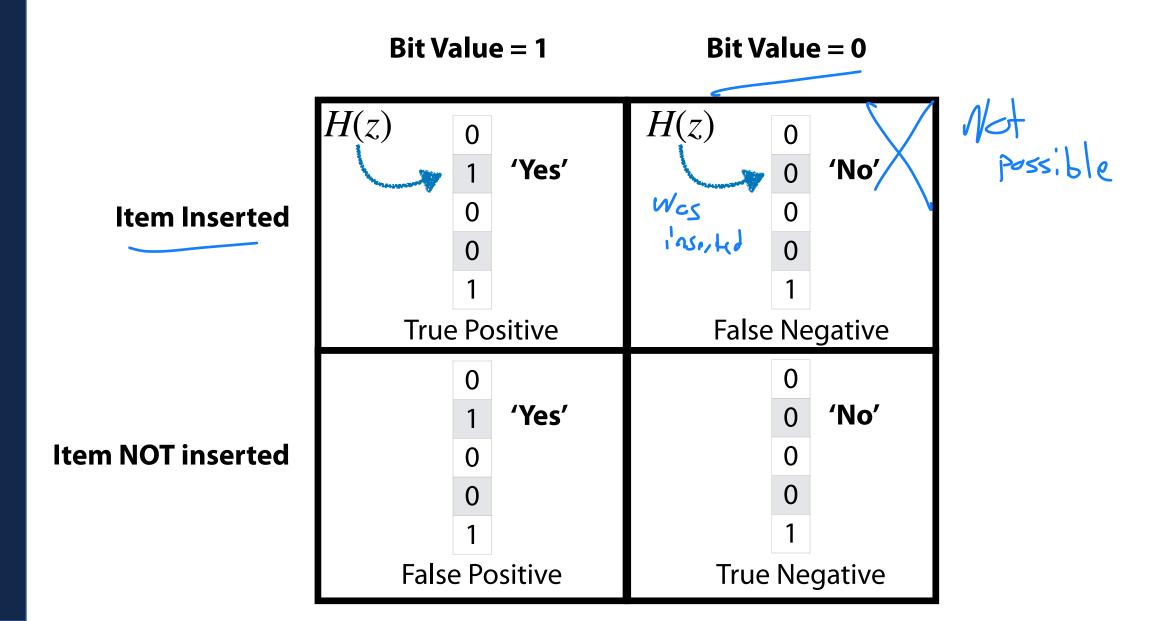
False Positive: Orade Says M

not m l actual m False Negative: not M

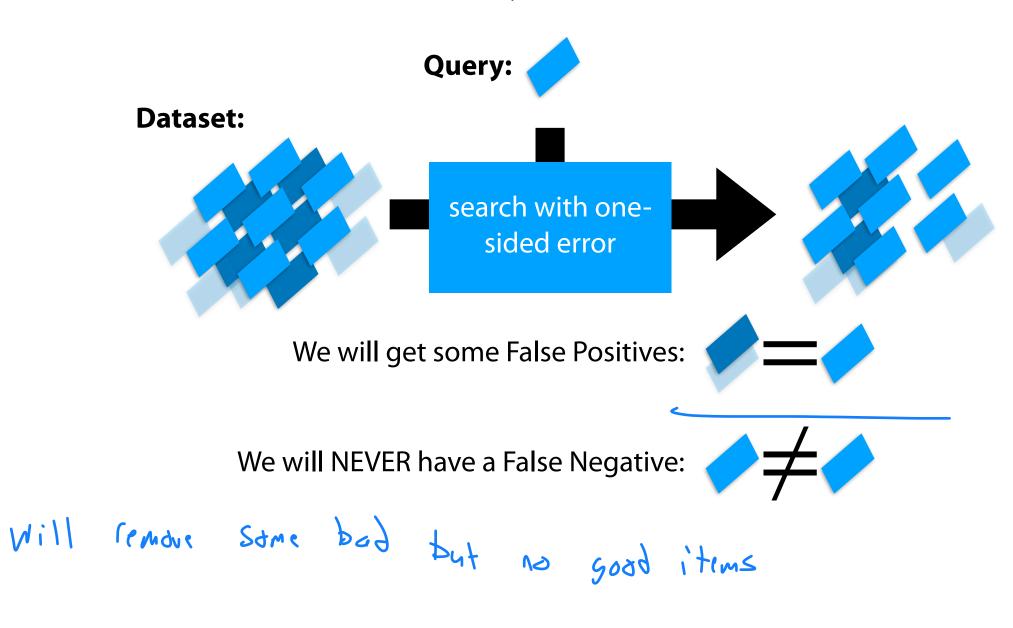
True Negative:

not m

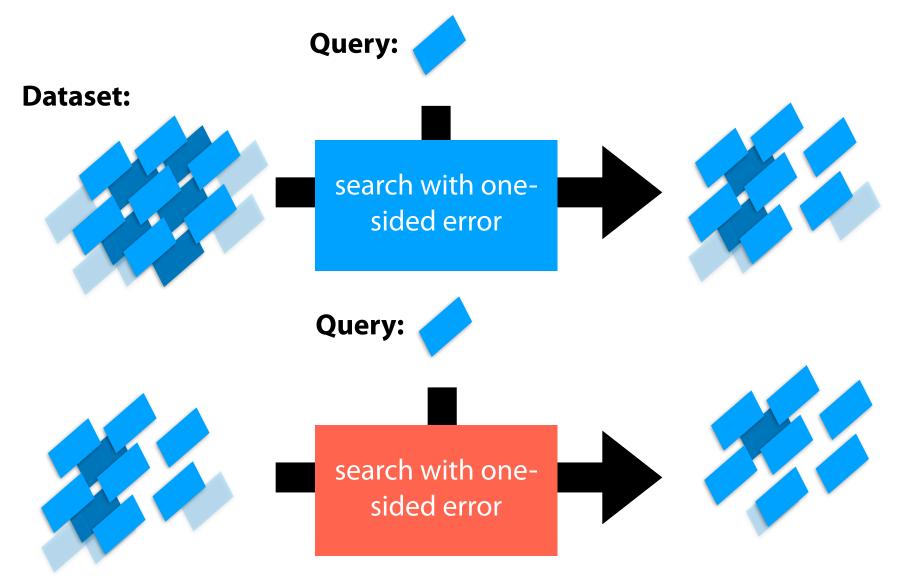
Imagine we have a **bloom filter** that **stores malicious sites...**

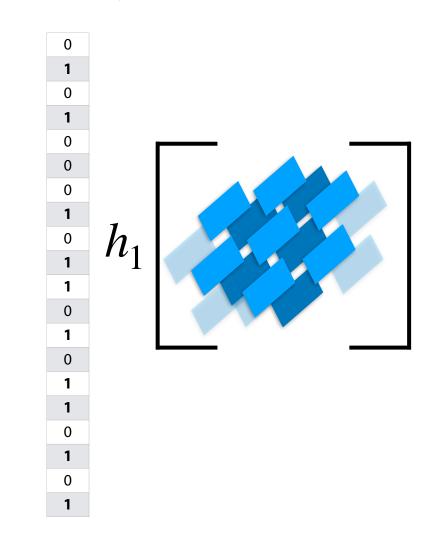


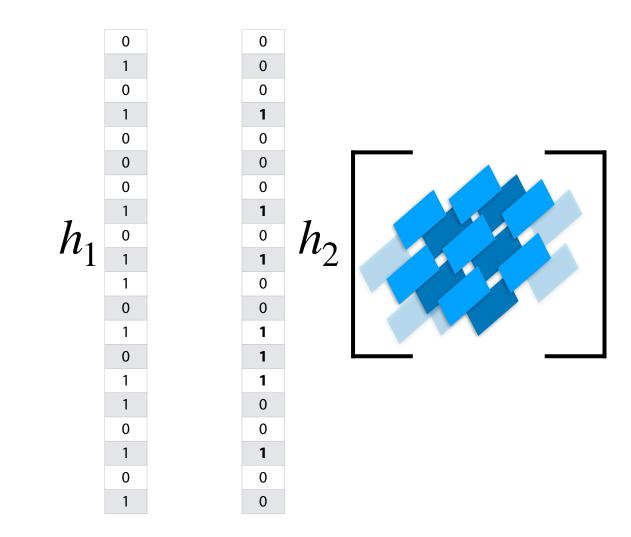
Probabilistic Accuracy: One-sided error

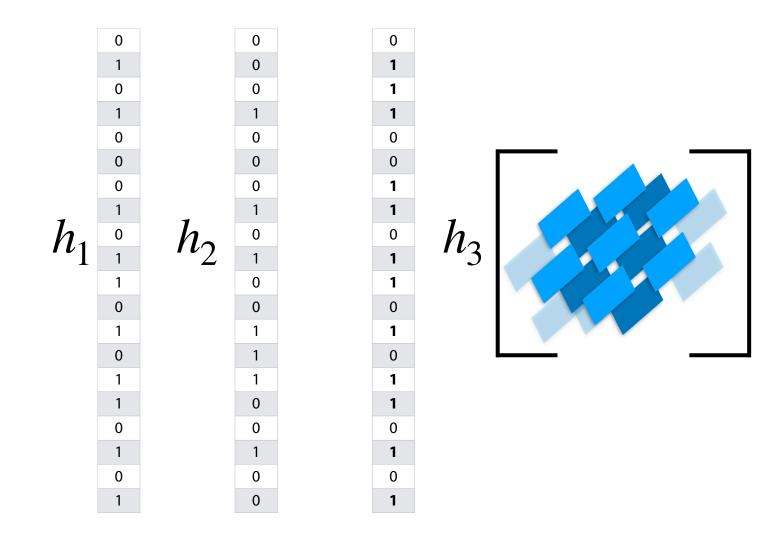


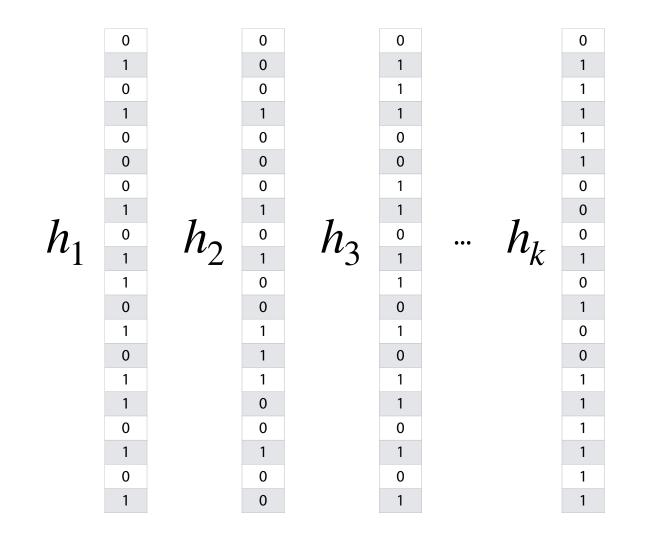
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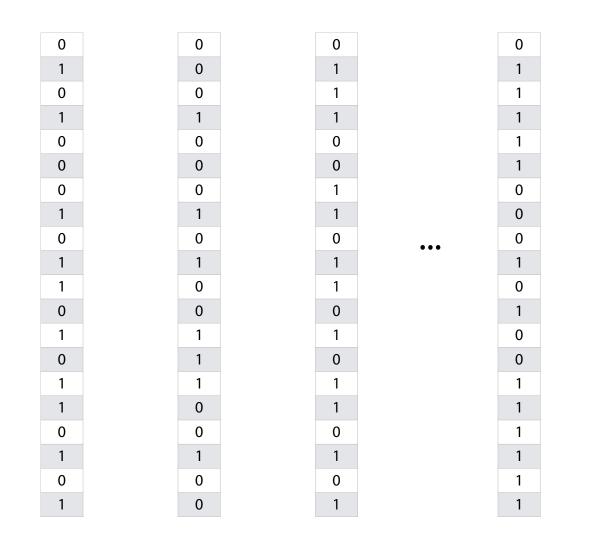




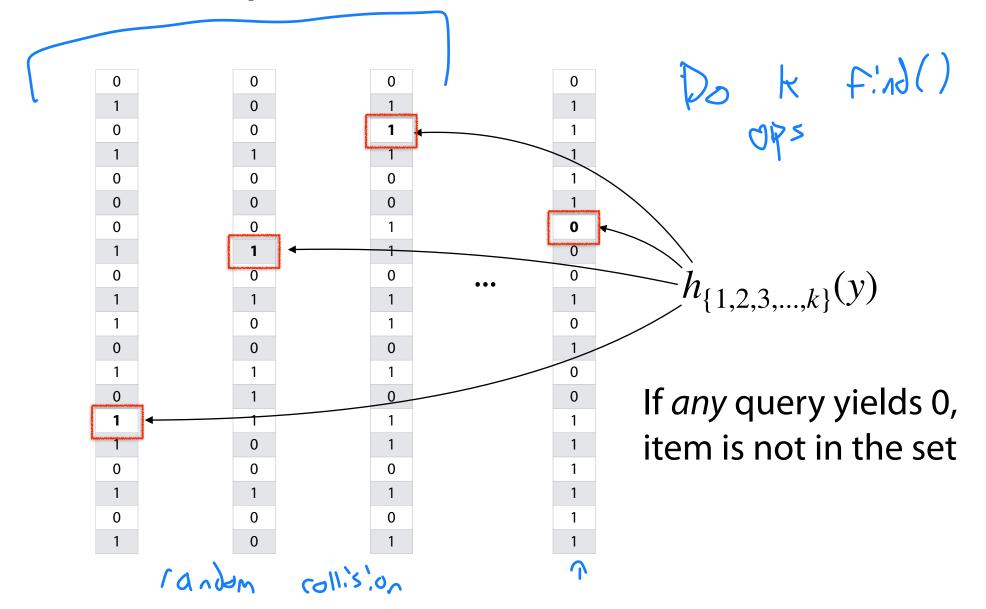


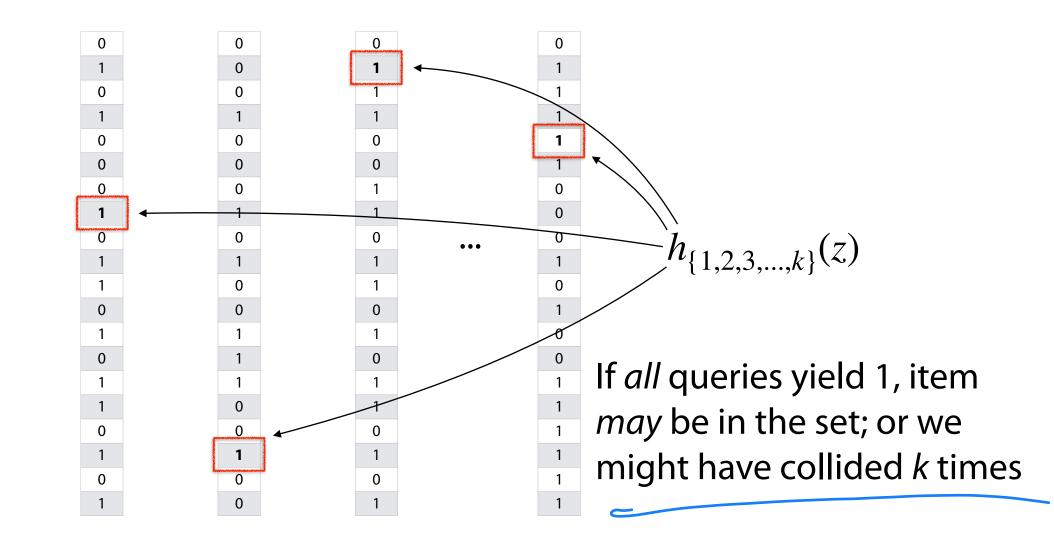






 $h_{\{1,2,3,\dots,k\}}(y)$





 $P^{K} \rightarrow P^{=0.5}$



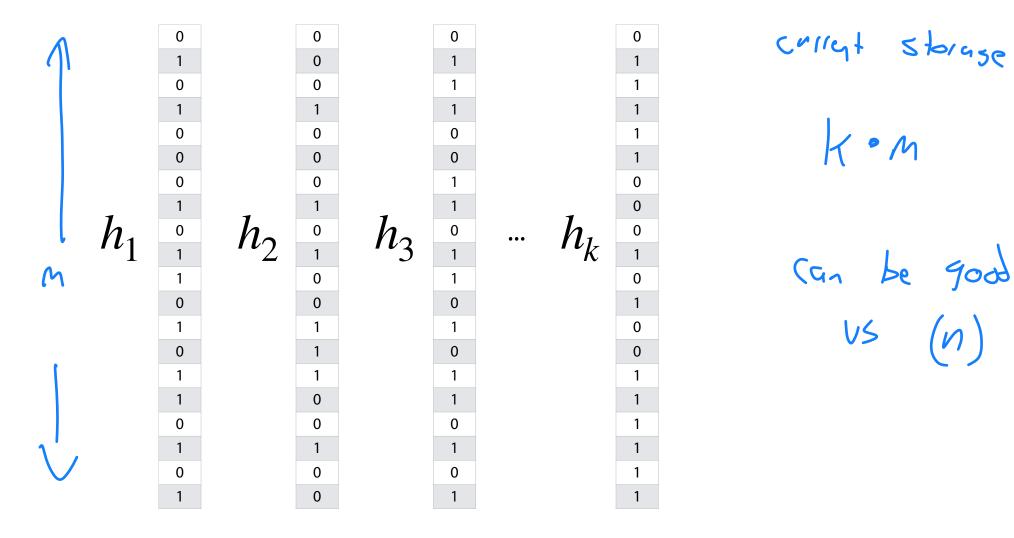
 $(0.5)^{10} \approx 0.00097$

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?

K · M



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

 $S = \{6, 8, 4\}$ c/ $h_1(x) = x \% 10$ $h_2(x) = 2x \% 10$ $h_3(x) = (5+3x) \% 10$ -> 6 -> 8 \rightarrow Y 1) Hash k times 2) Insert K times 2 6

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

 $h_2(x) = 2x \% 10$ $h_1(x) = x \% 10$ $h_3(x) = (5+3x) \% 10$ 1 Ż find(1) <u>~</u>2 -73 1) Hush # time/ 2) Do K lookurs (findli) > False 1 4 1 5 0 ~6 1 find(16) 6 1 7 Is true (this is fabre Positive,) 1 K 8 9 1

Bloom Filter

A probabilistic data structure storing a set of values

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

Insert / Find runs in: _

Delete is not possible (yet)!