Data Structures and Algorithms Bloom Filters 2

CS 225 Brad Solomon November 3, 2023



Department of Computer Science

Extra Credit Project Submissions

~110 teams submitted extra credit projects.

Drafted TAs to do a first pass grading of some of the major topics

Each TA-graded project is graded by two TAs for fairness

Mentors will (hopefully) be assigned sometime next week

Quick announcements on MPs

MP_Traversal had the lowest plagiarism rate of any assignment!

MP_mazes is due next week

The next MP will NOT be released next Monday

(> Aent nont malay?)

Quick announcements on Exams

Next exam is next Monday

Look at topic list / do practice exam

Make sure you thoroughly understand the coding question.



Discuss bit vector operations and potential extensions to bloom filters

Memory-Constrained Data Structures

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



Bloom Filter: Deletion

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



0



find(16)らり(16)= 2 Bit is one So Yes! _find(20) 6 (3 h(20) 7 This is (3 h(20) 7 This is (3 tes! a False positive

find(3) 4 h/3) 4 No Bloom Filter: Search Throw out information Trade off is loss in gaugery $H(\alpha)$ The bloom filter is a *probabilistic* data structure! Vale 5/0000 of the time, item is not present If the value in the BF is 0: $H(x_1)$ $H(\beta)$ L7No false regatives $H(x_2)$ If the value in the BF is 1: (> The object might be present $H(x_3)$ either I insert () $H(x_A)$ 0 I hash collided

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: Oracle says Malilions / Predikter Says Malilions False Positive: Orade says safe / Predicter Says Malicips False Negative: A ctail Malicians) Predict Safe Safe | predict Safe **True Negative:**



Probabilistic Accuracy: One-sided error



Probabilistic Accuracy: One-sided error





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



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



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



R J.ffernt hashes



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





FPR=L

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?

 $\begin{pmatrix} 1 \\ 5 \end{pmatrix}^{K} \begin{pmatrix} 1 \\ 3 \end{pmatrix}^{16} = 0.000976$ Power of repeated trials one-sided Prior

K=10



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

 $S = \{6, 8, 4\}$ 0 🔿 $h_1(x) = x \% 10$ $h_2(x) = 2x \% 10$ $h_3(x) = (5+3x) \% 10$ \mathcal{A} Я 6 8 Insert GII hashes glurays

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







Bloom Filter: Error Rate $h_{\{1,2,3,\ldots,k\}}$ Given bit vector of size *m* and 1 SUHA hash function What's the probability a specific bucket is 1 after After I insort Prod chases From 1/m one object is inserted? $P(burket=1) = \frac{1}{10}$ т Un next insect Same probability given k SUHA hash function? K Mot type! This is had to write!



Bloom Filter: Error Rate

Given bit vector of size *m* and *k* SUHA hash function

K buinness b/cK content trials $h_{\{1,2,3,\ldots,k\}}$

m

What's the probability a specific bucket is 1 after *n* objects are inserted?

 $-\left(\left| -\frac{1}{m} \right)^{\prime}\right)$







Bloom Filter: Optimal Error Rate

To build the optimal hash function, fix **m** and **n**!

Claim: The optimal hash function is when $k^* = \ln 2 \cdot \frac{m}{n}$

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

(2)
$$\frac{d}{dk} \left(1 - e^{\frac{-nk}{m}}\right)^k \approx \frac{d}{dk} \left(k \ln(1 - e^{\frac{-nk}{m}})\right)$$

Bloom Filter: Optimal Error Rate

Claim 1:
$$\left(1 - \left(1 - \frac{1}{m}\right)^{nk}\right)^k \approx \left(1 - e^{\frac{-nk}{m}}\right)^k$$

$$\left(1-\frac{1}{m}\right)^{nk} = e^{\ln\left[\left(1-\frac{1}{m}\right)^{nk}\right]}$$

$$= e^{\ln\left[\left(1-\frac{1}{m}\right)\right]nk}$$

$$pprox e^{\frac{-nk}{m}}$$

Bloom Filter: Optimal Error Rate

Claim 2:
$$\frac{d}{dk} \left(1 - e^{\frac{-nk}{m}}\right)^k \approx \frac{d}{dk} \left(k \ln(1 - e^{\frac{-nk}{m}})\right)$$

Fact:
$$\frac{d}{dx} \ln f(x) = \frac{1}{f(x)} \frac{df(x)}{dx}$$

TL;DR:
$$min [f(x)] = min [ln f(x)]$$

Derivative is zero when
$$k^* = \ln 2 \cdot \frac{m}{n}$$



Bloom Filter: Optimal Parameters -> Lob. Bloom

 $k^* = \ln 2 \cdot \frac{m}{n}$

Given any two values, we can optimize the third

$$n = 100$$
 items $k = 3$ hashes $m =$

$$m = 100$$
 bits $n = 20$ items $k =$

$$m = 100$$
 bits $k = 2$ items $n =$

Bloom Filter: Optimal Parameters

$$m = \frac{nk}{\ln 2} \approx 1.44 \cdot nk$$

Optimal hash function is still O(m)!



n = 250,000 files vs ~10¹⁵ nucleotides vs 260 TB



n = 60 billion — 130 trillion

Bloom Filter: Website Caching





Maggs, Bruce M., and Ramesh K. Sitaraman. Algorithmic nuggets in content delivery. ACM SIGCOMM Computer Communication Review 45.3 (2015): 52-66.

Bitwise Operators in C++

Traditionally, bit vectors are read from RIGHT to LEFT

Warning: Vector<bool> doesn't do this but actual bits do!





Bit Vectors: Unioning

Bit Vectors can be trivially merged using bit-wise union.



Bit Vectors: Intersection

Bit Vectors can be trivially merged using bit-wise intersection.



Bit Vector Merging

What is the conceptual meaning behind **union** and **intersection**?

Imagine we have a large collection of text...



And our goal is to search these files for a query of interest...









	SRA	FASTA.gz	SBT
Leaves	4966 GB	2692 GB	63 GB
Full Tree	-	-	200 GB

Solomon, Brad, and Carl Kingsford. "Fast search of thousands of short-read sequencing experiments." *Nature biotechnology* 34.3 (2016): 300-302.

Solomon, Brad, and Carl Kingsford. "Improved search of large transcriptomic sequencing databases using split sequence bloom trees." *International Conference on Research in Computational Molecular Biology*. Springer, Cham, 2017.

Sun, Chen, et al. "Allsome sequence bloom trees." *International Conference on Research in Computational Molecular Biology*. Springer, Cham, 2017.

Harris, Robert S., and Paul Medvedev. "Improved representation of sequence bloom trees." *Bioinformatics* 36.3 (2020): 721-727.

Bloom Filters: Tip of the Iceberg



Cohen, Saar, and Yossi Matias. "Spectral bloom filters." *Proceedings of the 2003 ACM SIGMOD international conference on Management of data*. 2003.

Fan, Bin, et al. "Cuckoo filter: Practically better than bloom." *Proceedings of the 10th ACM International on Conference on emerging Networking Experiments and Technologies*. 2014.

Nayak, Sabuzima, and Ripon Patgiri. "countBF: A General-purpose High Accuracy and Space Efficient Counting Bloom Filter." 2021 17th International Conference on Network and Service Management (CNSM). IEEE, 2021.

Mitzenmacher, Michael. "Compressed bloom filters." *IEEE/ACM transactions on networking* 10.5 (2002): 604-612.

Crainiceanu, Adina, and Daniel Lemire. "Bloofi: Multidimensional bloom filters." Information Systems 54 (2015): 311-324.

Chazelle, Bernard, et al. "The bloomier filter: an efficient data structure for static support lookup tables." *Proceedings of the fifteenth annual ACM-SIAM symposium on Discrete algorithms*. 2004.

There are many more than shown here...