Randomized Project - Bloom Filter

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Outline

- Introduction
- Related Work
- Results
- 4 Applications

Introduction

- Enhancing Collaborative Spam Detection with Bloom Filters
 - Jeff Yan & Pook Leong Cho (Newcastle University)
 - IEEE 2006 22nd Computer Security Applications Conference

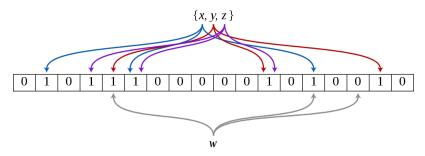
Spam Detection:

- Statistical vs. Signature-based Collaborative (SCSD)
- Using Bloom filters with SCSD:
 - O(1) look-ups
 - no huge database required



Bloom Filters

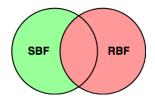
- Array of m bits (all initially set to zero)
- ullet k independent hash functions map to $\{0,...,m-1\}$
- Element x is inserted by setting locations $h_1(x), ..., h_k(x)$ to one



- Can introduce false positives
- $h_1(x) = ... = h_k(x) = 1$, when actually x was never inserted

Results of the Paper (Razor)

- Can basically use normal bloom filters, but...
- Razor supports deletion, which requires 2 Bloom filters:
 - Spam Bloom Filter (SBF), Revocation Bloom filter (RBF)



This achieves constant time look-up

Storing *n* signatures of length 160 bits:

- Bloom filters: 2m bits needed
- Hash table: at least 160 * n bits

Results of the Paper (DCC)

- Spam is "bulk". Count each time an email is reported
- Threshold value t (e.g. t = 20) indicates spam
- Signature x has count values

$$c[h_1(x)], c[h_2(x)], ..., c[h_k(x)]$$

- $\min\{c[h_1(x)], ..., c[h_k(x)]\}$ is an upper bound on the actual number of times x was inserted
- Storage:
 - Bloom filters: m * sizeof (cell) bits (e.g. 5m bits)
 - Hash table: at least n * size of(signature) + n * size of(count) bits bits

Results of the Paper (DCC)

Further heuristics when inserting signature x to decrease the False-Positive rate:

- **1** Only increment counters that equal $\min\{c[h_1(x)], ..., c[h_k(x)]\}.$
 - Global coincidental hits
- ② If two or more $h_i(x)$ map to the same cell, only increment that cell once.
 - Local coincidental hits

Variables

- *n* Number of elements/emails
- m Filter size, i.e, the number of bits allowed each encoded email vector
- k Number of hash functions constructed

Simulation and Results

They ran simulations to determine how much the 2 heuristics improved the counting Bloom filter

- Instead of actual email signatures, they used random numbers
- 8 experiments of varying insertion sequences for n = 10,000 distinct elements, e.g.

$$\underbrace{x_1, x_2, ..., x_{10,000}}_{Round_1}, \underbrace{x_1, x_2, ..., x_{10,000}}_{Round_{20}} \underbrace{x_1, ..., x_1}_{20}, \underbrace{x_2, ..., x_2}_{20}, \dots ..., \underbrace{x_{10,000}, ..., x_{10,000}}_{20}$$

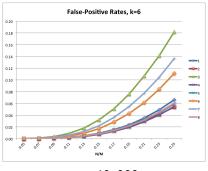
• Configurations for $k = \{4, 6, 8\}$ and $m = \{80K, 160K, 320K\}$

Findings: Heuristics significantly lowered False-Positive rates. Frequency of insertion, as well as order, also affect False-Positive rates.

What happens as n gets increasingly large?

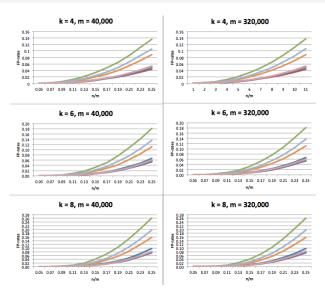
At what ratio of n/m do False-Positive rates become undesirable? Experiment:

- $k = \{4, 6, 8\}$ $m = \{40K, 80K, 160K, 320K\}$
- $n/m = \{0.05, 0.07, ..., 0.25\}$



$$m = 40,000$$

Experiments



- Each colored line represents an experiment
- Tested by varying both k and m
- As n/m or k increases, false positives increase

Applications

- Not necessarily great for spam detection...
- Useful when querying a large database, where looking up/reading data on disk is expensive.
 - Can have a Bloom filter overhead that is first queried to see if the item you are searching for even exists in the database
 - If the Bloom filter says it doesn't exist, then you know you don't have to do an expensive look-up, otherwise, the data you are looking for most likely exists
- Determining whether an email is in your contact list shorter strings than full emails

Future Work

- Test experiments on actual spam detection data
 - Spambase https://archive.ics.uci.edu/ml/datasets/Spambase
 - TREC 2007 Public Corpus https://plg.uwaterloo.ca/cgi-bin/cgiwrap/gvcormac/foo07
 - Enron http://www2.aueb.gr/users/ion/data/enron-spam/
 - SMS Spam http://www.esp.uem.es/jmgomez/smsspamcorpus/, https://www.kaggle.com/uciml/ sms-spam-collection-dataset#spam.csv

Conclusion

- Discussed Bloom Filters
- Discussed Related Papers
- Described Simulation Setup
- Described Experiments and Results
- Discussed Applications & Future Work

Questions?

