# Identifying Frequent Flows in Large Traffic Sets through Probabilistic Bloom Filters

 Yanjun Yao, Sisi Xiong, Jilong Liao, Michael Berry, Hairong Qi, and Qing Cao Department of Electrical Engineering and Computer Science University of Tennessee, Knoxville, TN, US
 Email: {yyao9, sxiong, jliao2, mberry, hqi, cao}@utk.edu

Abstract—In many network applications, accurate traffic measurement is critical for bandwidth management and detecting security threats such as DoS (Denial of Service) attacks. In such cases, traffic is usually modeled as a collection of flows, which are identified based on certain features such as IP address pairs. One central problem is to identify those "heavy hitter" flows, which account for a large percentage of total traffic, e.g., at least 0.1% of the link capacity. However, the challenge for this goal is that keeping an individual counter for each flow is too slow, costly, and non-scalable. In this paper, we describe a novel data structure called the Probabilistic Bloom Filter (PBF), which extends the classical bloom filter into the probabilistic direction, so that it can effectively identify heavy hitters. We analyze the performance, tradeoffs, and capacity of this data structure, as well as developing two extensions to improve its accuracy and flexibility. Our study also investigates how to calibrate this data structure's parameters, where we prove our developed method achieves the Nash Equilibrium using game theory. We use real network traces collected on a web query server and a backbone router to test the performance of the PBF, and demonstrate that this method can accurately keep track of all objects' frequencies, including websites and flows, so that heavy hitters can be identified with constant time computational complexity and low memory overhead.

*Index Terms*—Network Measurement, Traffic Analysis, Data Structures, Bloom Filter.

# I. INTRODUCTION

In managing today's complex Internet backbones, accurate traffic monitoring and measurements are crucial for many applications, including short-term purposes such as security needs (e.g., detecting traffic hot-spots, intrusions, and cyberattacks) and long-term traffic engineering purposes (e.g., rerouting common traffic, expanding the capacity for frequently chosen links) [1], [2], [3]. One central problem in such applications is to identify heavy hitters, i.e., those most frequent flows, by keeping track of flow frequencies based on real-time traffic. Given that the number of flows between commercial end host pairs can be extremely large [1], [4], however, keeping a counter for each flow usually requires more memory than available on limited hardware resources, such as routers. Existing methods have addressed this problem through sampling and counting, such as NetFlow [5], where one of Npackets is sampled and counted. However, such methods can only sample a small portion of the entire traffic, leading to inaccurate results and over- or under-estimates.

Given such challenges, in this paper, we address the problem on how we can efficiently construct estimates for the frequencies of all traffic flows so that heavy hitters can be identified. To this end, we aim to build an approximate histogram of all traffic flows with limited memory space, so that we can easily identify whether a flow is "heavy" when it is encountered in the ongoing traffic. The key assumption in our approach is that, for traffic management purposes, approximate knowledge on flows' frequency is already sufficient to identify heavy flows, as long as the frequency estimates can provide reliable upper and lower bounds associated with their most likely values. On the other hand, for some flows, if we need to obtain their accurate frequencies, we can add a few extra counters to serve such needs separately.

Our work in this paper is enabled by extending a compact, hashing-based data structure called the *bloom filter*. A Bloom Filter (BF) is a data structure that is designed to answer a query on whether an element exists in a set. Its basic idea is to hash an element to k different locations in a bit array, and sets these locations to all 1s when inserting this element to the set. Being a randomized method, it allows for false positives, but the space savings often outweigh its drawbacks. BFs were originally introduced for database applications, but recently they have received great attention also in the networking area (see [6], [7] as two surveys).

Based on the bloom filter, we investigate how to store frequency estimations, rather than set memberships. Note that previous work has addressed the "accurate" version of this problem by proposing counting bloom filters (CBF) [8], [6], [7], where the bit vector is replaced by a counter vector. The cost is that the CBF design usually consumes memory space that is one order of magnitude higher than the original BF. In this paper, as we are concerned with the constrained memory of devices such as routers, our extension is still based on bit vectors rather than counter vectors. Specifically, we present a probabilistic version of the bloom filter and its operations, so that we can provide estimates of flow frequencies using a small amount of memory, based on which we can provide reliable identification of heavy hitter flows. Formally, we define the problem as follows: given a multi-set S, we would like to identify those items that appear for more than f times. Note that items may be provided in a stream, as is the case of traffic flows, where IP addresses in headers are used to denote flows.

The central idea of our design, by extending the classical bloom filter, is it performs probabilistic counting operations. Therefore, we call this new data structure as the *Probabilistic Bloom Filter* (PBF). The key difference is that whenever an

item is inserted, instead of flipping the hash locations from 0 to 1, we flip them with a pre-set probability of p. Such a paradigm shift does not need any extra memory space compared to the standard bloom filter. Hence, it is still highly compact and feasible to implement on memory-constrained devices. We then model the performance of the PBF rigorously through probabilistic analysis, and we outline our major contributions as follows:

- We present the Probabilistic Bloom Filter, which allows non-deterministic queries on item existence and frequency in data sets. We provide the PBF's APIs and demonstrate how they can be used by applications.
- We quantitatively study the performance of the PBF through analytical approaches, where we derive closedform results regarding its capacity, overhead, and performance.
- We extend the PBF into two variants: a counting PBF (C-PBF) and a time-decaying PBF (T-PBF), for additional application needs.
- We study the parameter selection of the PBF and its two variants, and we develop an algorithm that proves to achieve the Nash Equilibrium using game theory.
- We evaluate the performance of the PBF with realistic Internet traffic datasets collected from a web query dataset and a backbone router for one hour of time to demonstrate its effectiveness.

We emphasize that our approach is fundamentally probabilistic and approximate. In fact, it cannot always return accurate estimations, and may, if poorly designed, fail to identify heavy hitters in three ways: it can miss some large flows, it can wrongly insert some small flows to the report, or it may give an inaccurate estimate of some large flows. We call these three types of errors: false negatives, false positives, and counting errors. We demonstrate the intricate tradeoffs between performance and overhead: the higher the requirement of accuracy, the more memory overhead the design will incur in practice. Our work also quantifies such tradeoffs through theoretical analysis.

While our evaluations are based on network-related datasets such as web query logs and traffic traces, the methods in this paper should be general enough to be applied to several domains. Indeed, counting the number of events based on their types in large-scale datasets is a widely used building block for data analysis in computational sciences. For example, the recent discovery of the Higgs boson using the LHC [9], [10] draws conclusions based on statistical analysis for collected particle collision events. Properly configured versions of PBFs can be applied for such purposes when TB scale of streaming data need to be analyzed in nearly real time under computational and memory constraints.

The remainder of this paper is organized as follows: We survey related work in Section II. The problem formulation and design are described in Section III. The extensions of the PBF are presented in Section IV, and the performance evaluation is given in Section VI. We provide conclusions in Section VII.

# II. RELATED WORK

In this section, we describe the related work in three parts: first, the original Bloom Filter design, then, its variants, and finally, recent progress on traffic flow sampling and counting in Internet routers.

The bloom filter, which is proposed by Burton H. Bloom in 1970 [11], is a space-efficient randomized data structure that answers the question on whether an element is in a set. There are two basic operations: insert and query. Its space efficiency is achieved at the cost of false positives (an element is claimed to be inside a set when it is not). The accuracy of a bloom filter depends on the filter size m, the number of hash functions k, and the number of inserted elements n. The false negatives (an element is reported as not in a set when it is) never happens. Although originally conceived for database applications, the bloom filter recently has also received great attention in the networking area [6], [7], [12], [13], [14], [15], [16], [17], [18].

In its initial design, the BF did not address the issues of element duplicates, as it only considers simple sets. No matter how many times an item appears in a set, it is counted only once in the constructed BF. Counting Bloom Filters (CBFs) [8], [6], [7] have been designed to address this issue. They are based on the same idea as BFs, but they adopted fixed size counters (also called bins) instead of single bits in its vector design. When an item is inserted, the corresponding counters are increased, hence, duplicate information is maintained rather than lost. However, CBFs are different from our approach in that they are fundamentally deterministic, as they keep accurate counts of the number of duplicates for an item. Therefore, CBFs require memory overhead that is usually an order of magnitude higher than common BFs, which makes them less scalable to many flows. Another related work to ours, proposed by Shen and others [19], [20], developed the idea of the Decaying Bloom Filter, which extended the Counting Bloom Filter to support the removal of stale elements when new elements are inserted. However, our design does not use as much memory as the decaying bloom filter for processing the same amount of data. Finally, Kumar and others [21] proposed the Space-Code Bloom Filter (SCBF) (later extended to a multi-resolution version called the MRSCBF), which used a filter made up of a fixed number of groups of hash functions. During the insertion operation, one group of hash functions was randomly chosen for the element. For query, the number of groups containing the element was counted to estimate the frequency. However, as only open formulas were provided, estimating frequency was done by looking up a pre-computed table, which was very computationally intensive to be built. Such efforts differ in our work in that we present closedform results on modeling the performance of the proposed data structures. In our evaluation, we compare with both the CBF and the MRSCBF as they are the state-of-the-art baselines.

In recent years, the bloom filter has been widely used in network measurements [1]. Estan and others [22] applied Counting Bloom Filters to traffic measurement problems inside routers. The approach was based on the simple idea that if the counter for a flow increases beyond a threshold, it should be considered as a frequent flow. Zhao and others [23] used distributed Bloom Filters to find local icebergs (items whose frequency is larger than a given threshold), and then estimated global icebergs in a central server. Finally, Liu and others [24] proposed the Reversible MultiLayer Hashed Counting Bloom Filter(RML-HCBF), whose hash functions select a set of consecutive bits from the original strings as hash values, so that it may find elephant flows (large and continuous flow) using the counter values and thresholds. In contrast, the PBF we propose is based on approximate counting methods rather than accurate ones, thus saving on the memory overhead and processing speed.

# III. PROBABILISTIC BLOOM FILTER

In this section, we describe the design of the probabilistic bloom filter (PBF). We first present its programming interfaces, followed by the operations between multiple PBFs, and finally, an analysis of its properties, capacity, and performance.

A. Programming Interfaces

Algorithm 1 The PBF Insert Algorithm

1: p	rocedure INSERT(x)	▷ Insert operation
2:	for $j = 1 \rightarrow k$ do	
3:	$i \leftarrow h_j(x)$	
4:	$random_i \leftarrow Uniform(0,1)$	
5:	if $random_i < p$ then	
6:	$B_i \leftarrow 1$	
7:	end if	
8:	end for	
9: <b>e</b>	nd procedure	

Algo	rithm	2	The	PBF	Frequenc	y Quer	y Algorithm
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1: **procedure** FREQUENCY(x)▷ Frequency test operation  $counter \leftarrow 0$ for  $j = 1 \rightarrow k$  do 2: 3:  $i \leftarrow h_i(x)$ if  $B_i == 1$  then 4: counter + +5. 6: end if end for 7:  $f \leftarrow estimation(counter)$ 8: Q٠ return f 10: end procedure

The PBF provides two programming APIs, an insert operation and a frequency query operation, as illustrated in Algorithm 1 and Algorithm 2. For the insert operation, the primary change compared to a conventional bloom filter is that it uses a parameter p to decide whether to flip a bit from 0 to 1, when items are inserted. Note that as an optimization, we do not need to read the bit's value before we set it to 1, thereby reducing the number of memory accesses for the insert operation. For the frequency query algorithm, it adds up the number of 1s in the k bits as determined by the hashing functions, and uses statistical inference methods to obtain an approximate frequency of the data item in the data set. We will describe the details of this estimation operation in Section III-C.

TABLE I Symbols Used in Analysis

f	The frequency threshold of heavy-hitter items
k	The number of hashing functions
m	The length of the bit-vector
n	The total number of flows or items in a dataset
p	The probability for setting a bit to 1
y	The expected number of 1s in the bit-vector
$\hat{y}$	The observed number of 1s in the bit-vector
$\theta$	The probability that a bit has been set to 1

### B. Properties of the PBF Operations

There are several operations for multiple PBFs, including the union and halving of PBFs, which are facilitated by the bit-vector nature of PBFs. Given two multi-sets  $S_1$  and  $S_2$ , suppose that they are represented by two PBFs, B1 and B2. We can calculate the PBF that represents the union set  $S = S_1 \cup S_2$  by taking the OR of their PBFs:  $B = B1 \vee B2$ assuming that the bit vector length m and the hash functions are identical. The merged filter B represents the aggregate frequency of an item belonging to  $S_1$  or  $S_2$  as belonging to the set S.

The second operation is halving. If the PBF size m is divisible by 2, halving allows us to store the original multiset in a shorter bit vector. This can be achieved by bit-wise ORing the first and second halves of the PBF's bit vector together. To insert or query the new PBF, the range of the hashing functions also needs to be updated by applying the mod(m/2) operation to their outputs.

# C. Performance Modeling of PBFs

Because the PBF introduces one additional parameter p, it has different properties compared to the original BF. In this section, we model the performance of the PBF by studying the relationship between the frequency of items and the number of bits that are flipped in the bit vector. Table I shows the notations we use in the following analysis.

We first consider what happens if there is just one item. We assume that this item has hashed positions that are distributed uniformly, as is true for the hashing functions we choose in our implementation. The probability that it flips a particular bit is p/m. Therefore, for a given bit, the probability that it is not set to 1 is given by

$$1 - \frac{p}{m}.$$
 (1)

With the PBF, there are k hashing functions for inserting any item. Hence, the probability that none of them will set a specific bit to 1 is given by

$$(1-\frac{p}{m})^k.$$
 (2)

After inserting n items into the bloom filter, the probability that a given bit is still zero is going to be

$$P(n,0) = (1 - \frac{p}{m})^{kn}.$$
(3)

Thus, the probability of a bit being set to 1 is

$$P(n,1) = 1 - (1 - \frac{p}{m})^{kn}.$$
(4)

The expected number of 1s of these k bits, denoted as g(p, m, n, k), is

$$g(p,m,n,k) = (1 - (1 - \frac{p}{m})^{kn}) * k \approx (1 - e^{-\frac{kpn}{m}}) * k.$$
(5)

Note that this approximation for the expected value is true only when p/m is sufficiently small. This constraint is true because our picked m is usually large, and p is usually much smaller than 1. Observe that P(n, 1) exists for every bit regardless of what items are inserted. Therefore, this value corresponds to a "background noise", which means that some bits will be set due to other items being inserted. Notice that when m increases, the background noise will decrease. If n increases, the noise increases. To obtain the frequency estimation of items, we have to take this noise into account.

Next, for a certain item that appears f times, it will invoke the *insert* API f times. Therefore, the probability for any of the k bits mapped by hash functions to still be zero is

$$P(f, 0) = (1 - p)^f \approx e^{-pf}$$
, and

the probability of this bit to be 1 is

$$P(f,1) = 1 - (1-p)^{f} \approx 1 - e^{-pf}.$$
(7)

(6)

Again, we assume that  $p \ll 1$ , which is true for our selection of p values. Clearly, whether a bit is set to 1 is determined by two factors: the probability that it is set to 1 by an item's insert operation as illustrated by P(f, 1), and the probability of the background noise as illustrated by P(n-f, 1) (in Eq. 4). These two factors are independent of each other. Therefore, the probability for an element to remain 0 as (since neither the background noise nor the repeated items have set it to 1) is given by

$$P(n-f,0) \times P(f,0) \approx e^{-\frac{pk(n-f)}{m}} \times e^{-pf}.$$
 (8)

Next, as each bit can be considered as a Bernoulli experiment, its "success" probability  $\theta$  can be considered as the event that a bit has been set as 1. Here we denote that

$$\theta = 1 - P(n - f, 0) \times P(f, 0) \tag{9}$$

Therefore, denote the total number of 1s as Y, we have that

$$Y|\theta \sim Bin(k,\theta). \tag{10}$$

Therefore, we know that  $E(Y|\theta) = k\theta$  and  $V(Y|\theta) = k\theta(1-\theta)$ . If we denote the expected number of bits that are set to 1 in the k mapped bits for an element in the PBF as y, we have

$$y = (1 - P(n - f, 0) \times P(f, 0)) \times k = k(1 - e^{-\frac{pk(n - f)}{m}} \times e^{-pf}).$$
(11)

Based on the observation results for the proportion of bits that have been set, we can denote its value as  $\hat{y}$ , and define  $\hat{\theta} = \hat{y}/k$ . We can use  $\hat{\theta}$  as an unbiased estimator of  $\theta$ , and since  $\theta$  is derived based on the frequency f, we can then estimate f as follows (by solving the Equation 11 for f)

$$f = \frac{knp + m\ln\left[1 - \frac{y}{k}\right]}{(k-m)p}.$$
(12)

Next, we calculate the confidence interval for f, by approximating it using a normal distribution based on the central limit theorem. This is also the so-called Wald Method, whose

formula for confidence interval is given by

$$\hat{\theta} \pm z_{\frac{1}{2}\alpha} \sqrt{\frac{1}{n} \hat{\theta} \left(1 - \hat{\theta}\right)} \tag{13}$$

where  $\hat{\theta}$  is the proportion of successes, and  $z_{\frac{1}{2}\alpha}$  is the critical z value with a tail area of  $\frac{1}{2}\alpha$  of the standard normal curve. Based on this formula, we can derive the lower and upper bounds for f as shown below:

$$f_{min} = \frac{knp + m\ln\left[\frac{k-\hat{y}}{k} + \sqrt{\frac{\left(1 - \frac{k-\hat{y}}{k}\right)(k-\hat{y})}{k^2}} z_{\frac{1}{2}\alpha}\right]}{(k-m)p}$$
(14)

$$f_{max} = \frac{knp + m\ln\left[\frac{k-\hat{y}}{k} - \sqrt{\frac{\left(1 - \frac{k-\hat{y}}{k}\right)(k-\hat{y})}{k^2}} z_{\frac{1}{2}\alpha}\right]}{(k-m)p}$$
(15)

As an illustrative example, suppose we want to filter data traffic with 100K flows, where frequent flows are defined as those with a frequency to be at least one percent of the total, i.e., 1K flows. We pick a bloom filter size of m = 2M bits  $(1M = 10^6)$ , and let k = 1000. We select p as 0.0006 (we will explain this later in parameter selection). In this case, the frequent flow is expected to have 467 bits in 1000 bits set as 1. Conversely, if indeed, 467 bits are set, the estimated number of flows is 999, with a 95% confidence interval as [905, 1098]. On the other hand, if somehow the value of  $\mu - 2\sigma = 435$ bits are set, the number of estimated flows is 902, with a 95% confidence interval as [813, 995]. Note that, however, this interval does not contain 1000 as it is a 95% confidence interval. One may want to use the 99% confidence interval to capture a larger range. In this case, the 99% interval gives the bound as [797, 1013], which indeed contains the 1000 in its range.

#### D. Selection of Parameters for PBFs

One critical challenge in using the PBF for analyzing datasets is that it needs to set several parameters properly, such as m, k, and p. Choosing such parameters improperly will reduce its capabilities and increase errors. Further, due to the probabilistic nature of the PBF, its parameters need to be set differently compared to conventional bloom filters. Therefore, in this section, we study how to set the parameters for the PBF, and define its capacity.

**Problem Formulation:** Given a known number of items n and the threshold for frequent flows f, how do we choose m, k, and p properly? Similarly, given m, k, and p, how do we estimate the PBF's capacity to handle large n and f?

To answer this question, we first find the constraints for m, k, and p, and try to optimize the model performance by minimizing m and k, which correspond to the memory overhead (m) and computational overhead (k).

The first constraint for choosing the right parameters is to limit the background noise as shown in Equation 5. As m increases, the background noise will decrease, assuming a fixed k. Therefore, to keep the noise below a certain threshold  $\epsilon$ , we require:

$$g(p, m, n, k)/k \le \epsilon,$$



Fig. 1. Relation between p and the estimation Fig. 2. bounds.

so that

$$m \ge \frac{-knp}{\log(1-\epsilon)}.\tag{16}$$

Note that we assume that  $\epsilon$  is chosen as an appropriately low threshold, e.g., 0.1.

The second constraint concerns the estimation accuracy as shown in Equation 12. To ensure the estimation accuracy, the observed value of  $\hat{y}$  should not be too small or too large compared to k. Since Equation 11 gives the expected value of y, we require that the ratio between y and k should lie between  $\epsilon$  and  $1 - \epsilon$ , therefore,

$$\epsilon \le \frac{k(1 - e^{-\frac{pk(n-f)}{m}} \times e^{-pf})}{k} \le 1 - \epsilon.$$
(17)

Based on this result, we can then obtain the estimation range of f, which is denoted by the lowest f value and the highest f value that can be accurately estimated, as a function of k, m, and p as

$$f_{lowerbound} = \frac{knp + \log(1 - \epsilon)m}{(k - m)p},$$
(18)

$$f_{upperbound} = \frac{knp + \log(\epsilon)m}{(k-m)p},$$
(19)

also, observe that the real value of f should be located within this estimation range, we have that

$$f_{lowerbound} \le f \le f_{upperbound}.$$
 (20)

This formula, therefore, gives the third constraint. To illustrate the meanings of these constraints, especially on how they affect the choices of p, we consider the following way to illustrate possible choices of parameter values. To minimize m, we simply set m as the lower bound, using Equation 16, i.e.,  $m = \frac{-knp}{\log(1-\epsilon)}$ . Then we can establish the constraints for p as

$$-\frac{\log(1-\epsilon)}{n} \le p \le \frac{(n-f)\log(1-\epsilon) - n\log(\epsilon)}{nf}.$$
 (21)

On the other hand, the value of k can be chosen based on two considerations. First, if k is too large, it will incur too much computational overhead. Second, the value of k will affect the confidence interval calculated in Equations 14 and 15. The reason is that different values of k will lead to different lower and upper bounds, and their difference will be varying. To illustrate this, assume that we have chosen p as the upper bound, and m as the lower bound, and we set  $\epsilon = 0.1$ , we can calculate the ratio between the confidence interval of f to the estimated f as follows

$$Ratio = \frac{f_{max} - f_{min}}{f}$$

Assuming  $f \leq \epsilon n$ , meaning that the frequency under





Relation between k and the estimation Fig. 3. Relation between p and  $\theta$ .

estimation is not too large compared to the total number of items, n, we can find that

Ratio 
$$\approx 0.46 \ln \left( 0.1 + 0.59 \sqrt{\frac{1}{k}} \right) - 0.46 \ln \left( 0.1 - 0.59 \sqrt{\frac{1}{k}} \right).$$

We can verify that under this setting, this ratio decreases monotonically with k increases. In particular, if we let  $Ratio \leq 1/2$ , we can find that  $k \geq 141$ . Therefore, we suggest k must be chosen not lower than 150 to ensure good performance. Note that this value of k only incurs moderate computation overhead. Recent research on fast string hashing algorithms [25] has demonstrated optimized hashing functions can achieve a hashing throughput of a fraction of a CPU cycle per byte. Therefore, we can use a k value of larger than 1000 on multi-core workstations without having performance bottlenecks, as hashing a 100 byte string (e.g., a packet header) 1000 times only takes less than 0.1ms.

So far, we have outlined the way we should select parameters. To summarize, the procedure is as follows:

Algorithm 3 The PBF Parameter Selection Algorithm						
1:	<b>procedure</b> SELECT( <i>n</i> , <i>f</i> )	$\triangleright n$ and $f$ as known				
2:	Calculate the bounds for $p$ ,					
	and use its upper bound i	if possible (Equation 21)				
3:	Select a modest value for $k$	(assuming $k \ge 150$ )				
4:	Calculate the lower bound for	or $m$ (Equation 16)				
5:	<b>return</b> $p$ , $k$ , and $m$					
6:	end procedure					

Note that we require p to be chosen as its upper bound because a larger p will increase the accuracy of estimation, as long as this p value does not go beyond its upper bound. We now use a numerical example to illustrate possible choices for p, k, and m. Assume that n = 100K and f = 1000. If  $\epsilon = 0.1$ , according to Equation 21, we have  $1.05 \times 10^{-6} \le p \le 0.0022$ . We now evaluate the effects of p. By keeping m bounded according to Equation 16 and k = 2000 (k can also take any other constant value), we change the value of p and study its effects on the confidence interval. The results are shown in Figure 1. As illustrated, a larger p indeed leads to better confidence intervals, as long as the value of p does not go over its upper bound. In fact, the narrowest confidence interval in Figure 1 is precisely when p = 0.0022. After p goes over this upper bound (specified by Equation 21), the confidence interval becomes larger again. The underlying reason for this interesting phenomenon is that when p is too large, it will over-saturates the bit vector too early, hence impairing the performance of estimations.

Next we investigate the effects of m. Suppose that we have chosen a moderate value of p for general cases, where p is chosen as 0.0006. We hope to see how k can affect confidence intervals, by changing k from 200 to 8000. As shown in Figure 2, a larger k will lead to better confidence intervals, at the cost of more computational overhead.

Finally, we illustrate the effects of p on  $\theta$ , which is the percentage of the 1s in the k bits after all flows are inserted. Figure 3 shows the results. Observe that as long as p is chosen according to Equation 21,  $\theta$  will be bounded by  $[\epsilon, 1-\epsilon]$ , which is expected. On the other hand, if p is larger than the upper bound,  $\theta$  will be over-saturated; if p is lower than the lower bound,  $\theta$  will be under-saturated.

## E. The Maximum Estimating Frequency

Once the PBF parameters are chosen, given properly chosen p, k, and m, one critical problem is its estimation errors. Given that our goal of this paper is to identify heavy-hitters, we are most concerned about the errors that are caused by oversaturation of bit vectors, i.e., when the frequency of the flow is too high. Motivated by this observation, we next define the "Maximum Estimating Frequency (MEF)", which is defined as follows:

**Maximum Estimating Frequency (MEF):** Given a set of PBF parameters, what is the maximum value of f that it can still estimate accurately, where the k bit vector is at most saturated for  $1 - \epsilon$ ?

To derive MEF, we use Equation 19, and replace m with the result from Equation 16. Therefore, the maximum value that f can reach is given by

$$f_{MEF} = \frac{n(\log(1-\epsilon) - \log(\epsilon))}{np + \log(1-\epsilon)}.$$
(22)

In reality, for specific PBF settings, the value of p is only a single value. Hence, if we set p as a constant, for a large n, we then have

$$\lim_{n \to +\infty} f_{MEF} = \frac{\log(1 - \epsilon) - \log(\epsilon)}{p}.$$

Hence, this approximation can be used to estimate the real MEF that a PBF setting can handle. If  $\epsilon = 0.1$ , the capacity is roughly 2.2/p. For example, if p = 0.001, this PBF can count up to cardinality of around 2,200. For all items with a frequency above this threshold, the PBF can still report that they are *at least* 2,200, but their real value is not reported due to bit vector saturations.

#### IV. EXTENSIONS OF THE PBF

In this section, we describe two extensions of the PBF for potential applications: a counting PBF (C-PBF) and a timedecaying PBF (T-PBF).

# A. C-PBF: Counting PBF Design

In this section, we introduce the C-PBF, a counting variant of the PBF that extends its capability with more memory usage. The idea of the C-PBF is simple: it replaces each bit in the PBF with a w-bit counter. Whenever an item is inserted, instead of deciding on whether flipping one bit from 0 to 1, it will determine with a probability of p whether to increase this w-bit counter. Formally, this updated algorithm is shown in Algorithm 4.

Algorithm	4	C-PBF	Insert	A	lgorithm
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1:	<b>procedure</b> INSERT( <i>x</i> )	▷ Insert operation
2:	for $j = 1 \rightarrow k$ do	
3:	$i \leftarrow h_j(x)$	
4:	$Counter_i \leftarrow B[i]$	
5:	$random_i \leftarrow Uniform(0,1)$	
6:	if $random_i < p$ then	
7:	$Counter_i = Counter_i + 1$	
8:	end if	
9:	end for	
10:	end procedure	

We can now derive the performance of the C-PBF. Observe that for each counter, it may be updated by two sources: the background noise caused by other items and the insertion operations of the frequent flows. Therefore, if we denote these two sources with two random variables  $X_1$  and  $X_2$ , we have

$$X_1 \sim Bin(k(n-f), p/m), \tag{23}$$

$$X_2 \sim Bin(f, p),\tag{24}$$

where  $X = X_1 + X_2$ . In this scenario, we can use the Poisson distribution to approximate these two distributions, so that the sum of the distributions will remain Poisson. In that case, the size of counter is bounded as  $\max_{i=1}^{m} \log_2(X_i)$ . Based on this result, considering that we observe k counters (assuming k hashing functions), we can denote their values as  $x_i$ , where  $i \in [1, k]$ . We can then use the average of these observations on  $x_i$  to estimate the MLE of f as

$$\hat{f} = \frac{knp - m\hat{x}}{(k-m)p}, \text{ where } \hat{x} = \sum_{i=1}^{n} x_i/n.$$
(25)

### B. Selection of Parameters for C-PBF

Although C-PBF uses counters for predicting the element frequencies, it is still critical to choose proper ranges for m, k, and p to increase prediction capabilities and reduce errors. The problem we need to solve is formulated as follows: given a known number of items and the threshold for frequent flows f, how do we choose m, k, and p properly? The same as what we discussed in PBF, to answer this question, we also need to find the constraints for m, k, and p to optimize its performance by minimizing m and k to achieve minimal memory overhead (m) and computational overhead (k).

Similar to PBF, limiting the background noise is the first constraint for choosing the right parameters. For a fixed k, the background noise will decrease, while m increases. In that case, to keep the background noise below a certain threshold  $\epsilon'$ , we need

$$\frac{Bin(kn, p/m)}{2^s} < \epsilon', \tag{26}$$

so that

$$m > \frac{knp}{2^s \epsilon'}.\tag{27}$$

On the other hand, as the memory is limited in networking devices, the size of the C-PBF is bounded by the memory capacity c and the counter size s. For a fixed counter size s, the size of m should be bounded as

$$\frac{knp}{2^s\epsilon'} < m \le \frac{c}{s}.\tag{28}$$

Secondly, to ensure the prediction accuracy, we should not under-saturate or over-saturate the counter. Therefore, we require that its ratio of the maximum counter value should lie between  $\epsilon'$  and  $1 - \epsilon'$ , so we have

$$\epsilon' \le \frac{kn\frac{p}{m} + fp}{2^s} \le 1 - \epsilon'.$$
<sup>(29)</sup>

Assume that  $\epsilon'$  is chosen as an appropriately low threshold, e.g., 0.1, with the second constraint, we can obtain the prediction range of f as a function of k, m, and p as:

$$f_{lowerbound} = \frac{knp - m2^s \epsilon'}{(k-m)p},\tag{30}$$

$$f_{upperbound} = \frac{knp - m2^s(1 - \epsilon')}{(k - m)p}.$$
(31)

Next, if we assume that the real value of f should be located within this prediction range, we have

$$f_{lowerbound} \le f \le f_{upperbound}.$$
 (32)

In that case, when we minimize m and set up  $m > \frac{knp}{2^s \epsilon'}$ , we can establish the constraints for p as:

$$\frac{2^{s}\epsilon'}{n} (33)$$

Finally, the value of k needs to be carefully selected. If k is too large, it would incur too much computation overhead, so is the background noise, as the probability of different elements mapping to the same filter bucket would be large. However, if k is too small, the deviation of counter values would be large. Now, assume that  $p = \frac{f2^s \epsilon' + n2^s - 2n2^s \epsilon'}{fn}$ ,  $m = \frac{knp}{2^s \epsilon'}$ , and  $\epsilon' = 0.1$ , then the ratio between the confidence interval of f to the predicted f is

$$Ratio = \frac{f_{max} - f_{min}}{f}.$$
(34)

Assume  $f \leq n\epsilon'$ , as we know that the counter value is following the Poisson Distribution with a parameter of  $\lambda$ , where the  $E(X) = Var(X) = \lambda$ , then

$$X_{max} = (\lambda + 1)\left(1 - \frac{1}{9(\lambda + 1)} + \frac{1.96}{3\sqrt{\lambda + 1}}\right)^3,$$
 (35)

$$X_{min} = \lambda (1 - \frac{1}{9\lambda} - \frac{1.96}{3\sqrt{\lambda}})^3,$$
 (36)

then

$$f_{max} = \frac{2^{s} \epsilon' * n - X_{max} n}{2^{s} \epsilon' - n \frac{f 2^{s} \epsilon' + n 2^{s} - 2n 2^{s} \epsilon'}{f n}},$$
(37)

$$f_{min} = \frac{2^{s} \epsilon' * n - X_{min} n}{2^{s} \epsilon' - n \frac{f 2^{s} \epsilon' + n 2^{s} - 2n 2^{s} \epsilon'}{f n}}.$$
 (38)

In that case, we can find that

$$Ratio = \frac{X_{min} - X_{max}}{2^s \epsilon' - X},$$
(39)

as

$$\lambda = fp + \frac{k(n-f)p}{m} = 2^s (1-\epsilon'), \tag{40}$$

Then

$$Ratio = -1.25\left(\frac{0.9\left(1 - \frac{0.123457}{2^s} - \frac{0.688674}{2^{\frac{5}{2}}}\right)^3 2^s}{2^s} - \frac{(1 + 0.9 * 2^s)\left(1 + \frac{0.653333}{\sqrt{1 + 0.9*2^s}} - \frac{0.123457}{1.11111 + 2^s}\right)^3}{2^s}\right).$$
 (41)

Interestingly, observe from the above result, we observe that the performance metric *Ratio* is not affected by k, i.e., they are independent. In that case, we choose  $k = \frac{m}{n} \ln 2$ , which is the same as what has been done in the classical BF.

#### C. T-PBF: Time-decaying PBF Design

In this section, we introduce the second variant, the T-PBF, a time-decaying variant of the PBF that allows it to *forget* those frequent items that appeared in the distant past. The idea of the T-PBF is to introduce a new operation, decaying, which flips bits from 1 to 0 with a probability q over each time epoch T. This can be considered as an approximation for the *delete* operation. Formally, this updated algorithm is shown in Algorithm 5.

Alg	orithm 5 T-PBF Decaying A	lgorithm
1:	<b>procedure</b> DECAY( <i>x</i> )	▷ Decay operation
2:	for Every $T$ seconds do	
3:	for $j = 1 \rightarrow k$ do	
4:	$i \leftarrow h_j(x)$	
5:	if then $B[i] == 1$	l
6:	$random_i \leftarrow U$	Uniform(0,1)
7:	if $random_i <$	q then
8:	$B[i] \leftarrow 0$	
9:	end if	
10:	end if	
11:	end for	
12:	end for	
13:	end procedure	

We now demonstrate the long term behavior of the T-PBF. For simplicity, we consider operations in epochs, and the decaying operation only occurs at the end of each epoch.

Observe that for each bit, for each epoch, it may either start with bit 1 or 0. If it starts with 1, it may be flipped to 0 at the end of the epoch with a probability of q. However, if it starts with 0, it may be flipped to 1 first with a probability shown in Equation 11, and then flipped back to 0 with a probability of q. Therefore, we can use a discrete time Markov chain to describe these operations. In particular, because the transitions exhibit different probabilities at the beginning and the end of each epoch, we can model it with a time-inhomogeneous chain with a seasonal variation. In this case, we have that the seasonal period d = 2, and the transition probability as the following (assuming  $n \ge 0$ , while 2n and 2n+1 stand for the index of epochs)

$$P(2n) = {\begin{smallmatrix} 0 \\ 1 \end{smallmatrix}} \left[ \begin{array}{cc} 0 & 1 \\ 1-\alpha & \alpha \\ 0 & 1 \end{array} \right], \quad P(2n+1) = {\begin{smallmatrix} 0 \\ 1 \end{smallmatrix}} \left[ \begin{array}{cc} 0 & 1 \\ 1 & 0 \\ \beta & 1-\beta \end{array} \right],$$

where we have:

$$\alpha = 1 - e^{-\frac{pk(n-f)}{m}} \times e^{-pf}, \text{and}$$
(42)

$$\beta = q. \tag{43}$$

To analyze this seasonal chain, we can add a supplementary variable to create a homogeneous chain. The new chain contains four states: A(0, a), B(0, b), C(1, a), D(1, b). Its transition matrix is shown below, followed by the corresponding Markov chain illustration.



Fig. 4. The Markov Transition for the T-PBF

This Markov chain has four communicating classes, A, B, C, and D. They are all recurrent classes, as well. In the long term, this chain has the following stationary distribution:

$$P(s) = \begin{cases} \frac{\beta}{2\alpha+2\beta} & \text{for } s \equiv A, \\ \frac{\beta}{2\alpha+2\beta} & \text{for } s \equiv B, \\ \frac{\alpha}{2\alpha+2\beta} & \text{for } s \equiv C, \\ \frac{\alpha}{2\alpha+2\beta} & \text{for } s \equiv D. \end{cases}$$

As expected, the results show that in the long term, if a flow is no longer available, f will decrease to 0, and its  $\alpha$  will quickly converge to  $1 - e^{-\frac{pkn}{m}}$ , which is smaller. Therefore, in the long run, most bits will be reset to 0 (as demonstrated by the increased probability of states A and B in Figure IV-C. This validates the design of the time-decaying nature of the T-PBF. On the other hand, if a flow re-appears with a large f, its transition will mostly stay in states C and D, which means that more bits will be set due to the flow existence. Note that the epoch of the T-PBF can also be dynamic: as the PBF will lose the ability of prediction when all k bits are set to 1s, when a large percentage of k bits for any element are set to 1, the decaying process can be triggered.

## D. Selection of Parameters for T-PBF

To perform accurate frequency estimations with T-PBF, it is critical to choose proper values for parameters, such as m, k, p, and q. As T-PBF extends PBF by introducing one additional parameter q, and the decaying operations are only triggered at the end of each period, we can use the same algorithms to choose the values of m, k, and p, based on what we discussed in Section III-D. As a result, in the following part, we will discuss how to choose the value for q.

For an element, at the end of the first epoch, the probability of a bit is still 1 is

$$\hat{\theta_1} = \theta_1 * (1 - q),$$
(44)

where *theta* is as defined in Equation 9.

Then at the end of the second epoch, the probability of a bit is 1 would be a joint effect of the first two epoch, therefore

$$\theta_{2}^{'} = (\theta_{2} + \theta_{1}^{'} - \theta_{2} * \theta_{1}^{'}) * (1 - q).$$
(45)

Finally, at the end of the ep-th epoch, the probability of a bit is 1 would be

$$\theta_{ep}^{'} = (\theta_{ep} + \sum_{i=1}^{ep-1} \theta_{i}^{'} - \theta_{ep} \prod_{i=1}^{ep-1} \theta_{i}^{'}) * (1-q), \qquad (46)$$

as in long term,  $\theta_{ep} \prod_{i=1}^{ep-1} \theta'_i \to 0$ , we can have the estimate probability as

$$\theta_{ep}^{\prime} \approx \sum_{i=1}^{ep} \theta_i (1-q)^{ep-i}.$$
(47)

To guarantee the prediction accuracy, we need to prevent over-saturating the k bits for any element. Therefore, we need to make sure that

$$\forall \sum_{i=1}^{ep} y_i (1-q)^{ep-i} \le k(1-\epsilon),$$
(48)

where ep is the number of passing epochs,  $\epsilon$  is a threshold with a small value, and  $y_i$  is the number of bits set to 1s with the insertion of an element in the *i*th epoch. As a result of above requirements, we can rewrite this equation as

$$\max \sum_{i=1}^{ep} y'_i (1-q)^{ep-i} \le k(1-\epsilon),$$
(49)

where  $y'_i$  is the number of 1 generated by the most frequent element to the bit array. By setting  $y'_i = y_{max} = \max_{i=1}^{ep} \max_{j=1}^{t} y_{ij}$ , which is the maximum number of 1s set by the element with the largest frequency in all epochs, so that t is the number of elements in each epoch, then

$$\sum_{i=1}^{ep} y_{max} (1-q)^{ep-i} \le k(1-\epsilon),$$
 (50)

then

$$(1 - e^{-\frac{pk(n-f_{max})}{m}}e^{-pf_{max}})\frac{1 - (1-q)^{ep}}{q} \le 1 - \epsilon.$$
 (51)

For a long run, where  $(1-q)^{ep} \to 0$ , then we have:

$$\frac{1 - e^{-\frac{pk(n-f_{max})}{m}}e^{-pf_{max}}}{1 - \epsilon} \le q \le 1.$$
(52)

# V. PBF PARAMETER SELECTIONS ARE NASH EQUILIBRIUMS

As discussed in the previous sections, the estimation accuracy of the PBF and the T-PBF would be bad if an improper combination of parameters is chosen and used. In the following, we demonstrate that our algorithms for parameter selection will always lead to Nash equilibriums in practice, i.e., further adjusting any parameter by itself will not lead to a better performance [26]. We do not, however, claim that the parameter selection is globally optimal, as we find such optimality may depend on the particular application requirements. Indeed, the tradeoffs between the parameters are complex, and optimizing for any single global metric out of context is not particularly meaningful. Our proof is based on modeling the rules for parameters as the strategies in a noncooperative game. Specifically, we prove that:

*Theorem 5.1:* The parameter selection algorithm for the PBF (Algorithm 3) outputs a set of parameters that forms a Nash equilibrium.

*Proof:* To prove that the PBF parameter selection algorithm leads to Nash equilibriums, we define a normal form game as a triple  $(N, (S_i)_{i \in N}, (c_i)_{i \in N})$ , where

- N is the set of players, and N = 3;
- $S_i$  is the set of strategies of the player *i*. Each player can choose to follow or not to follow a specific rule for a parameter (p, m, or k), as specified in Algorithm 3, where  $S_1 = \{Follow\_P, Not\_Follow\_P\}$ ,  $S_2 = \{Follow\_K, Not\_Follow\_K\}$ , and  $S_3 = \{Follow\_M, Not\_Follow\_M\}$ ;
- $S = S_1 \times S_2 \times S_3$  is the set of states;
- a state is  $s = (s_1, ..., s_3) \in S$ ;
- c<sub>i</sub>: S → ℜ is the cost function of player i ∈ N. In the state s player i has a cost of c<sub>i</sub>(s). The cost is defined as the difference between the upper and lower estimation bounds, which represents the estimation accuracy.

In general, the cost value is computed as (based on results from 18 and 19):

$$C(p,k,m) = f_{max} - f_{min} = \frac{m(\ln \epsilon - \ln(1-\epsilon))}{(k-m)p}.$$
 (53)

Based on the Nash Theorem [26], which claims that every finite normal form game has a mixed Nash equilibrium, the PBF parameter selection game has a mixed Nash equilibrium. In that case, in the following part, we will show that the state as generated in the PBF parameter selection algorithm forms one Nash equilibrium for the game.

As described in the game, each player has two strategies as following or not following the rule for a specific parameter. For player 1, the cost for the game is as shown in Table II. where FP is the abbreviation of  $Follow_P$ , and NFP represents  $Not_Follow_P$ . Other abbreviations follow the same format. Therefore, the cost for player 2 and player 3 are shown in Table III and Table IV.

TABLE II Cost for player 1.

	(FK,FM)	(NFK,FM)	(FK,NM)	(NK,NM)
FP	C(p, k, m)	C(p, k', m)	C(p, k, m')	C(p, k', m')
NFP	C(p', k, m)	C(p', k', m)	C(p', k, m')	C(p', k', m')

TABLE III Cost for player 2.

	(FP,FM)	(NFP,FM)	(FP,NFM)	(NFP,NFM)
FK	C(p, k, m)	C(p', k, m)	C(p, k, m')	C(p', k, m')
NFK	C(p, k', m)	C(p', k', m)	C(p, k', m')	C(p', k', m')

Now, we will find the best response strategy of the three players. Recall that the parameter selections can be summarized as  $-\frac{\log(1-\epsilon)}{n} \le p \le \frac{(n-f)\log(1-\epsilon)-n\log(\epsilon)}{nf}, k \ge 150$ , and  $m \ge \frac{-knp}{\log(1-\epsilon)}$ . Consequently, we can find the opposite of these strategies as  $p' < -\frac{\log(1-\epsilon)}{n}$  or  $p' > \frac{(n-f)\log(1-\epsilon)-n\log(\epsilon)}{nf}$ , k < 150, and  $m < \frac{-knp}{\log(1-\epsilon)}$ , as NFP, NFK, and NFM, respectively. Let us start from player 1, whose costs for choosing different strategies are as follows:

$$C_1(FP) = C(p, k, m) + C(p, k', m) + C(p, k, m') + C(p, k', m'),$$
(54)

$$C_1(NFP) = C(p', k, m) + C(p', k', m) + C(p', k, m') + C(p', k, m'),$$

$$+ C(p', k', m'),$$
(55)

When the k bits mapped by an element are saturated, we define the difference between upper and lower bounds of the estimated frequency as  $\infty$ , as the PBF can no longer provide an accurate estimation for the frequency. This corresponds to the case that we choose  $p' > \frac{(n-f)\log(1-\epsilon)-n\log(\epsilon)}{nf}$ , where we have  $C_1(NFP) = \infty$ , so  $C_1(NFP) > C_1(FP)$ . On the other hand, when  $p' < -\frac{\log(1-\epsilon)}{n}$ , we also have that  $C_1(NFP) > C_1(FP)$ , as the cost is monotonically increasing with the decreasing of p. Hence, the best response strategy for player 1 is *Follow\_P*.

Following the same procedure, the costs for player 2 under different strategies are:

$$C_1(FK) = C(p, k, m) + C(p', k, m) + C(p, k, m') + C(p', k, m'),$$
(56)

$$C_1(NFK) = C(p, k', m) + C(p', k', m) + C(p, k', m') + C(p', k', m').$$

$$+ C(p', k', m').$$
(57)

As the cost is monotonically increasing with the decreasing of k, and k > k', then we have that  $C_1(FK) < C_1(NFK)$ . In that case, the best response strategy for player 2 is  $Follow_K$ .

Now let us have a look at player 3, whose costs under different strategies are as follows:

$$C_1(NFK) = C(p, k', m) + C(p', k', m) + C(p, k', m') + C(p', k', m').$$
(58)

$$C_1(FM) = C(p, k, m) + C(p', k, m) + C(p, k', m) + C(p', k', m),$$
(59)

$$C_1(NFM) = C(p, k, m') + C(p', k, m') + C(p, k', m') + C(p', k', m').$$

$$+ C(p', k', m').$$
(60)



Fig. 5.Relation between p and Fig. 6.Relation between k and Fig. 7.Relation between m and Fig. 8.Relation between q andestimation bounds.estimation bounds.estimation bounds.estimation bounds.

TABLE IV Cost for player 3.

	(FP,FK)	(NFP,FK)	(FP,NFK)	(NFP,NFK)
FM	C(p,k,m)	C(p',k,m)	C(p,k',m)	C(p',k',m)
NFM	C(p,k,m')	C(p',k,m')	C(p,k',m')	C(p',k',m')

As the cost function can be rewritten as  $C(p, k, m) = \frac{\ln \epsilon - \ln(1-\epsilon)}{\frac{kp}{m} - p}$ , we observe that its value will increase with the decrease of m. As m > m', we have that  $C_1(NFM) > C_1(FM)$ . In that case, the best response strategy is  $Follow_M$  for player 3.

Above all, we have proven that  $\{Follow\_P, Follow\_K, Follow\_M\}$  is a Nash equilibrium for the game. In another word, the parameter selection algorithm of PBF is a Nash equilibrium.

Next, we consider the algorithm parameter selection for T-PBF, and we prove that:

*Theorem 5.2:* The parameter selection algorithm for the T-PBF with regard to q generates a Nash equilibrium.

*Proof:* The proof is similar to the previous proof, by analyzing the effects of q to the strategies. The detailed proof is omitted due to space.

# VI. EVALUATION

In this section, we evaluate the performance of the PBF and its extensions, the C-PBF and the T-PBF, using one web query dataset and one real Internet traffic dataset. We compare the performance of our proposed algorithms with the following three baselines, in terms of estimation accuracy and memory usage:

- Counting Bloom Filter (CBF) [8]: as a well-known extension of the Bloom Filter, CBF uses counters to replace bits in each filter bucket. If the frequency of an element grows large, CBF requires allocation of larger memory space to hold all counters.
- Multi-Resolution Space-Code Bloom Filter (MRSCBF) [21]: the second extension, MRSCBF, employs multiple filters operating at different resolutions, where the frequency estimation is performed by looking up this number in a pre-computed lookup table.
- Random Sampled Netflow (RSN) [5]: RSN processes only one randomly selected packet out of n sequential packets, and then estimates the frequency for each sampled element based on its individual counter.

We emphasize that our approach is fundamentally probabilistic and approximate. Therefore, it may not be able to return accurate estimations, and may fail to identify heavy hitters in three ways: it may give inaccurate estimates, it may wrongly insert some small flows to the report, and it may miss some large flows. We call these three types of errors: estimation errors, false positives, and false negatives. We define these metrics as following:

- Estimation error ratio: this metric is defined as the difference, in percentage, between the estimated frequency and the real frequency.
- False positive ratio: the false-positive ratio is defined as the percentage of falsely reported heavy-hitters, whose frequencies are actually below f, among all flows whose frequencies are lower than f.
- False negative ratio: the false-negative ratio is defined as the percentage of falsely un-reported flows, whose frequencies are actually above f, among all flows whose frequencies are above f.

# A. Dataset A: Web Query Log Analysis

Our first evaluation dataset is a public web query dataset [27], where web query logs with 20M web queries are collected from 650K users over a period of three months. Each web query contains the user ID, searching keyword(s), a timestamp, and the web link that this user picks. The daily workload pattern is plotted in Figure 9, which shows a seven-day repeating pattern on the number of queries per day. In the following, we will use the PBF and the C-PBF to detect popular websites and key words, and use T-PBF to show the trend of the daily query number for the website *http://www.google.com*.

Note that due to the limited size of the dataset, this task can be finished with any conventional method. However, our goal is to demonstrate that our proposed methods can achieve a better memory usage while introducing only limited errors. The limited size of this dataset allows us to evaluate the performance of these methods easily.

1) Detecting Popular Websites: The dataset includes a total of 378,087 websites selected by users, where around 0.45% of them have a frequency above 100. We define the popular websites as those that appear for more than 100 times.

To detect popular websites, we first feed the dataset to a PBF with k = 150, p = 0.001, and m = 6M, where the parameters are selected based on the method in Section III-D. Note that the total memory use is only 0.75MB (6M bits). Figure 10 compares the estimated frequency and the real frequency of the first 500 popular websites. Observe that the estimations are matching real frequencies closely, with an average estimation error computed using the formula  $\frac{f_{estimated} - f_{real}}{f_{real}}$  as 4.7%.



Fig. 17. Detect popular key word Fig. 18. TDetect popular key word Fig. 19. C-PBF for the web query with the CBF for the web query with the MRSCBF for the web query with the RSN for the web query with the dataset dataset. dataset.

Detect popular key word Fig. 20. Detect popular key word dataset

The primary source of inaccuracies comes from the randomness when flipping a bit from 0 to 1. Furthermore, note that Figure 10 shows the upper and lower bounds of f estimations, which have much larger errors compared to the estimated value of f (the estimate of f is calculated based on Equation 12).

We also test the same dataset with the C-PBF, and the performance is comparable. Specifically, for C-PBF, we set up k = 50, p = 0.03, m = 0.6M. The counter size of C-PBF is set to 10 bits, so that the total memory overhead is the same as PBF. The average estimation error of C-PBF is 4.9%, as shown in Figure 11. With this setting of C-PBF, the maximum frequency that can be estimated is 34,105, according to Equation 25, which is sufficient for this dataset.

Next, we compare the performance of PBF/C-PBF with the baselines. The first baseline is CBF, where we set it up with suitable parameters. We choose to set the counter size to be 16 bits to accommodate the most frequent flows. Compared to the PBF, the CBF consumes 16 times of the memory as the PBF and C-PBF under similar settings. The advantages of CBF lie in that it does not provide approximate answers, as demonstrated by the Figure 12, where no confidence intervals are plotted. The second baseline is MRSCBF, where we set it up with l = 32, r = 5, and  $m_i = \frac{k_i n l}{\ln 2}$ , where  $k = \{3, 4, 6, 6, 6\}$ as mentioned in [21]. As shown in Figure 13, the average estimation error of MRSCBF is 10.1%, which is twice of the error of PBF and C-PBF. On the other hand, besides a long and offline pre-computation phase to get the lookup table, the MRSCBF takes  $\sum_{i=1}^{5} m_i = 436, 371, 391$  bits memory space, which is 72.7 times of the memory as the PBF and C-PBF. The only advantage of MRSCBF is that once the lookup table is built, it takes a few CPU cycles to estimate the frequency. In PBF, however, a constant k hashing functions need to be calculated. The third baseline is RSN, where we randomly sample one packet out of ten packets. As a result of the low sample rate, RSN only takes 11,063,808 bits memory space, which is 1.8 times of the memory as the PBF and C-PBF. According to Figure 14, the average estimation error of RSN is 11.3%. The major sources of errors are the randomness of sampling, and frequency estimation based on the sample counters and the predefined sample rate. Reduced estimation errors can be achieved, at the cost of taking more memory space.

2) Detecting Popular Keywords: In this dataset, there are 580,392 different keywords used for searching websites, where around 1.69% of them have a frequency above 100. We define the popular key words as those that appear for more than 100 times.

To detect popular keywords, we first feed the dataset to a PBF with k = 150, p = 0.0005, and m = 8M, where the parameters are selected based on the method in Section III-D. Note that the total memory use is only 1MB (8M bits). Figure 16 compares the estimated frequency and the real frequency of the first 500 popular key words. In general, the estimations are matching real frequencies closely, with an average estimation error as 4.9%, where the primary source of inaccuracies is the randomness when flipping a bit from 0 to 1.

On the other hand, we also test the same dataset with the

C-PBF, and get a comparable performance. For C-PBF, we set up k = 50, p = 0.02, m = 0.8M. The counter size of C-PBF is set to 10 bits, such that the total memory overhead is the same as PBF. The average estimation error of C-PBF is 3.5%, as shown in Figure 17. As the maximum frequency that can be estimated is 34, 105, according to Equation 25, there is no missing point in the figure.

Next, we compare the performance of PBF/C-PBF with the baselines. Compare to the first baseline CBF, which is set up with suitable parameters. We choose to set the counter size to be 18 bits to accommodate the most frequent flows. Compared to the PBF, the CBF consumes 18 times of the memory as the PBF and C-PBF under similar settings, as demonstrated by the Figure 18. For the second baseline MRSCBF, we also set up l = 32, r = 5, and  $m_i = \frac{k_i n l}{\ln 2}$ , where  $k = \{3, 4, 6, 6, 6\}$ . As shown in Figure 19, the average estimation error of MRSCBF is 9.7%, which is twice of the error of PBF and C-PBF. On the other hand, besides a long and offline pre-computation phase, the MRSCBF takes  $\sum_{i=1}^{5} m_i = 669,862,928$  bits memory space, which is 83.7 times of the memory as the PBF and C-PBF. The only advantage of MRSCBF is that once the lookup table is built, it takes a few CPU cycles to estimate the frequency, which is smaller compared to a constant k hashing functions need to be calculated in PBF. For the third baseline RSN, we also choose the sample rate as 0.1. As a result of that, RSN takes 18, 487, 258 bits of memory space, which is 2.3 times of the memory as the PBF and C-PBF. On the other hand, as shown in Figure 20, the average estimation error of RSN is 14.2%, which is around three times of the error of PBF and C-PBF.

3) Frequency Trend of a Popular Website: For the longterm detection, we can use T-PBF to reveal the trend of frequency of a popular website. Here we choose Google as our observing target. For the T-PBF, we set up k = 150, p = 0.001, m = 6M, and q = 0.8. We trigger the decay operation every time we finish processing one day's data. As shown in Figure 15, in general, T-PBF can estimate the frequency trend of the Google website well. On the other hand, when sudden changes occur, T-PBF cannot adapt fast enough due to the existence of previous epochs' data, which take time to cause the filter contents to decay.

#### B. Dataset B: Network Measurement Dataset Analysis



Fig. 21. Dataset traffic pattern, gener-Fig. 22. Dataset flow frequency, genated based on [28] erated based on [28]

In our second case study, we use PBF to analyze heavyhitters from Internet traffic traces. Our evaluation dataset contains passive traffic traces from CAIDA's equinix-chicago and equinix-sanjose monitors on high-speed Internet backbone links [28]. The dataset spans one hour of activity. First, we analyze the general traffic pattern of the trace, by counting the total number of packets collected in each minute. The traffic patterns are shown in Figure 21. In the following, we will use the PBF and the C-PBF to detect the heavy hitter flows, and use the T-PBF to detect the long-term trends of flow frequencies.

To differentiate between flows, we use pairs of source and destination IP addresses as the key for each flow. We then count the frequency of each flow in the whole dataset, where the results are shown in Figure 22. Observe that most of the flows have a frequency of only once or twice, but a small number of flows exist with a large frequency. In our following experiments, we define a threshold of 1,000 for those heavy hitters. These flows account for 0.12% of the total 96,854,555 flows, where the maximum frequency is 32,404,064.

1) Detecting Heavy Hitter Flows: First, we evaluate the estimation accuracy of the PBF. We focus on the data traces of the first 5 minutes when we apply PBF and C-PBF in this study. Later, we use T-PBF to handle the entire hour of data. The reason is that networked devices such as routers will most likely process datasets in a streaming manner, exactly as what T-PBF does, instead of processing the dataset in one operation.

For the first 5 minutes, there are 187, 116, 831 packets, which belong to 15, 454, 076 different flows. The number of flows with a frequency larger than 1,000 is 13,569. As there are limited resources on networking devices, we set p = 0.00005, k = 4,000, and m = 800M bits, according to the method in Section III-D. The comparison between the real frequency and the estimated frequency of the first 500 frequent flows (sorted by time) are shown in Figure 23. Observe that the estimated frequency and the real frequency are almost perfect matching each other. The only exceptions are those cases when the PBF loses its estimation ability as it's k hash bits are saturated with 1s, leading it to miss certain data points. However, we argue that such missing points have no effects on the identification of heavy hitters, as these missing points correspond to heavy flows with a high certainty.

Next, we use the same first 5 minute data to evaluate the estimation accuracy of the C-PBF. To fit the C-PBF into networking devices, we set up the size of the counter to be 10 bits, so that m = 80M, k = 400, and p = 0.0015. As shown in Figure 24, the C-PBF can estimate the frequency of flows equally accurately compared to PBF. On average, the estimated frequency is 3.83% larger than the real frequency, caused by the inherent approximate nature of C-PBF counting designs. This accuracy is comparable to the original PBF design.

Next, we compare with the baselines including the CBF, the MRSCBF, and the RSN. For the CBF method, we set k = 4000 and m = 800M. As shown in Figure 25, the estimated frequencies are also close to the real frequencies, with an average error as 3.76%. However, to prevent counter overflows, we have to set each counter to occupy 22 bits, which means that in terms of memory overhead, the CBF takes 22 times of the memory compared to PBF and C-PBF to deliver accurate counting performance. For the second baseline MRSCBF method, we set l = 32, r = 6, and  $k = \{3, 4, 6, 6, 6, 6\}$  to accommodate the most frequent flows.





Fig. 23. Comparison between the real fre- Fig. 24. quency and the estimated frequency with the quency and the estimated frequency with the quency and the estimated frequency with the PBF C-PBF.



Comparison between the real fre- Fig. 25. Comparison between the real fre-CBF.



Comparison of the real heavy hitter

Comparison between the real fre- Fig. 27. Comparison between the real fre- Fig. 28. Fig. 26. quency and the estimated frequency with the quency and the estimated frequency with the ratio and the detected heavy hitter ratio of the MRSCBE RSN. T-PBF.

Log(f)



Fig. 29. The average error rate of the PBF with Fig. 30. The false positive ratio of the PBF. Fig. 31. The false negative ratio of the PBF. p = 0.0001, and different k and m values.

To reduce the computation time of the look-up table, we set the maximum estimated frequency as 10,000, which has no effects on detecting heavy hitter flows. This is why the estimated frequency is never larger than 10,000 in the Figure 26. Excluding the flows with frequencies more than 10,000, the average estimation error of MRSCBF is 9.3%, which is more than twice of the error of PBF and C-PBF. On the other hand, it needs  $\sum_{i=1}^{6} m_i = 22, 117, 154, 656$  bits of memory to hold this MRSCBF, which means that the MRSCBF takes 27.6 times of the memory compared to PBF and C-PBF to estimate flow frequencies. For the RSN method, we set sample rate as 0.1. We can observe from Figure 27 that the average estimation error is 4.61%. On the other hand, RSN takes 1,304,986,970bits of memory space for this large dataset, which is 1.63 times of the memory compared to PBF and C-PBF.

2) Long-term Heavy Flow Detection with T-PBF: We next investigate the performance of long-term flow detection with the T-PBF on the whole dataset. For the T-PBF, we set p = 0.00005, k = 4000, m = 800M, and q = 0.3. The decay operation is triggered after processing one minute's traffic data. We calculated the detected heavy hitter ratio (the number of detected heavy flows over the total number of flows), the falsepositive ratio, and the false-negative ratio for each period. For both the false-positive ratio and the false-negative ratio, the

threshold on frequency is set as 1000. As shown in Figure 28, the detected heavy hitter ratio on average is 6.48% larger than the real heavy hitter ratio. The source of this inaccuracy comes from the accumulation of flow frequencies in the previous periods. In general, the T-PBF works as expected, as we can observe that it triggers small estimation errors and low memory overhead.



Fig. 32. The memory overhead of the PBF with different k values.

3) Effects of Parameter Selections: We next perform experiments to evaluate the effects of k on estimation errors, the false-positive ratio, and the false-negative ratio. With a given k, the computational overhead and query delay is constant. In the experiments, we choose p to be 0.00005 and 0.0001, respectively, and change k from 200 to 2000 with a step size 50. The value of m is computed from Equation 16 with different k and p values. As shown in Figure 29, the estimation error decreases with the increasing k values. This observation is consistent with our theoretical evaluation shown in Figure 2.

On the other hand, as shown in Figure 30, the maximum false-positive ratio is 0.000009, which is acceptably small. The false negative ratio shown in Figure 31 has a maximum value of 0.03, which is also acceptably small. Note that the false negative ratio is larger than the false-positive ratio, because the number of heavy hitter flows is much smaller than the number of non-heavy-hitter flows. We also observe that by increasing k, the false-positive and false-negative ratios are decreasing. Finally, with the increasing of k and p, the needed memory size is increasing as shown in Figure 32. This explains the tradeoff between the error rates and the memory overhead: a larger memory overhead typically leads to a better performance.

## VII. CONCLUSION

In this paper, we develop the probabilistic bloom filter (PBF), which extends conventional bloom filters to perform probabilistic counting operations. We provide the PBF's APIs to demonstrate how they can be used by applications, and quantitatively investigate the performance of the PBF through analytical approaches. The derived closed-form results answer our questions regarding the capacity, accuracy, and parameter selection of the PBF. In particular, we demonstrate the parameters chosen by our algorithms form Nash equilibriums. Finally, we also extend the PBF into two variants: a counting PBF (C-PBF) and a time-decaying PBF (T-PBF), for additional application needs. Our evaluation results based on two realistic datasets show that this design outperforms the existing, state-of-the-art approaches.

To the best of our knowledge, our work in this paper is the first probabilistic bloom filter that is designed to count large volume of data with adjustable capacity and accuracy. We hope our work can stimulate future work in this direction, and provide a basis for investigations towards better methods based on probabilistic counting and bloom filters.

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