Invertible Bloom Lookup Tables

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Abstract

We present a version of the Bloom filter data structure that supports not only the insertion, deletion, and lookup of key-value pairs, but also allows a complete listing of its contents with high probability, as long the number of key-value pairs is below a designed threshold. Our structure allows the number of key-value pairs to greatly exceed this threshold during normal operation. Exceeding the threshold simply temporarily prevents content listing and reduces the probability of a successful lookup. If later entries are deleted to return the structure below the threshold, everything again functions appropriately. We also show that simple variations of our structure are robust to certain standard errors, such as the deletion of a key without a corresponding insertion or the insertion of two distinct values for a key. The properties of our structure make it suitable for several applications, including database and networking applications that we highlight.

1   Introduction

The Bloom filter data structure [5] is a well-known way of probabilistically supporting dynamic set membership queries and it has been used in a multitude of applications (e.g., see [8]). The key feature of a standard Bloom filter is the way it trades off query accuracy for space efficiency, by using a binary array T (initially all zeroes) and k random hash functions, $h_1, \ldots, h_k$, to represent a set S by assigning $T[h_i(x)] = 1$ for each $x \in S$. To check if $x \in S$ one can check that $T[h_i(x)] = 1$ for $1 \le i \le k$, with some chance of a false positive. This representation of S does not allow one to list out the contents of S given only T. This aspect of Bloom filters is sometimes viewed as a feature, in settings where some degree of privacy protection is desired (e.g., see [2, 3, 18, 27, 30]). Still, in many domains one would benefit from a similar set representation that would also allow listing out the set's contents [17].

In this paper, we are interested not in simply representing a set, but instead in methods for probabilistically representing a lookup table (that is, an associative memory) of key-value pairs. Unlike previous approaches (e.g., see [6, 9]), we specifically desire a data structure that supports the listing out of all of its key-value pairs. We refer to such a structure as an *invertible Bloom look-up table* (IBLT).

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1.1 Related Work

Our work can be seen as an extension of the invertible Bloom filter data structure of Eppstein and Goodrich [17], modified to store key-value pairs instead of only keys. Our analysis, however, supersedes the analysis of the previous paper in several respects, in terms of efficiency and tightness of the analysis. In particular, our analysis demonstrates the natural connection between these data structures and cores of random hypergraphs, similar to the connection found previously for cuckoo hashing and erasure-correcting codes (e.g., see [16, 23]). In addition, our IBLT supports some usage cases and applications (discussed later in this section) that are not supported by a standard invertible Bloom filter.

Given its volume, reviewing all previous work on Bloom filters is, not surprisingly, beyond the scope of this paper (e.g., see [7, 8, 33] for some excellent surveys). Nevertheless, two closely related works include Bloomier filters [9] and Approximate Concurrent State Machines (ACSMs) [6], which are structures to store and track key-value pairs. Song *et al.* [35] store elements in a Bloom-like hash table in a fashion that is somewhat similar to that of a cuckoo hash table [15, 26, 29]. Cohen and Matias [12] describe an extended Bloom filter that allows for multiplicity queries on multisets. While an IBLT is not intended for changing values, it can be used in such settings where a key-value pair can be explicitly deleted and a new value for the key re-inserted. Again, an IBLT has additional features, including listing and graceful handling of overflow and counting multiplicities, which make it useful for several applications where these other structures are insufficient.

Another similar structure is the recently developed counter braid architecture [22], which keeps an updatable count field for a set of flows in a compressed form, with the compressed form arising by a careful use of hashing and allowing reconstruction of the count for each flow. Unlike an IBLT, however, the flow list must be kept explicitly to read out the flow counts, and such lists do not allow for direct lookups of individual values. There are several other differences from our work due to their focus on counters, but perhaps most notable is their decoding algorithm, which utilizes belief propagation. Additional work in the area of approximate counting of a similar flavor but with very different goals from the IBLT includes the well-known CM-sketch [13] and recent work by Price [32].

1.2 Our Results

We present a deceptively simple variation of the Bloom filter data structure that is designed for key-value pairs and further avoids the limitation of previous structures (such as [6, 9]) that do not allow the listing of contents. As mentioned above, we call our structure an *invertible Bloom lookup table*, or IBLT for short. Our IBLT supports insertions, deletions, and lookups in $O(k)$ time, where k is the number of random hash functions used (which will typically be a constant in practice). Just as Bloom filters have false positives, our lookup operation works only with constant probability, although this probability can be made quite close to 1.

Our data structure also allows for a complete listing of the key-value pairs the structure contains, with high probability, whenever the current number, n, of such pairs lies below a certain threshold capacity, t, a parameter that is part of the structure's design. This listing takes $O(t)$ time. In addition, because the content-listing operation succeeds with high probability, one can also use it as a backup in the case that a standard lookup fails—namely, if a lookup fails, perform a listing of key-value pairs until one can retrieve a value for the desired key.

Our IBLT construction is also space-efficient, requiring space[1] at most linear in t, the threshold number of keys, even if the number, n, of stored key-value pairs grows well beyond t (for example, to polynomial in t) at points in time. One could of course instead keep an actual list of key-value pairs with linear space, but this would require space linear in n, i.e., the *maximum* number of keys, not the target number, t, of keys. Keeping a list also necessarily requires more computationally expensive lookup operations than our approach supports.

We further show that with some additional checksums we can tolerate various natural errors in the system. For example, we can cope with key-value pairs being deleted without first being inserted, or keys being inserted with the same value multiple times, or keys mistakenly being inserted with multiple values simultaneously. Interestingly, together with its contents-listing ability, this error tolerance leads to a number of applications of the IBLT, which we discuss next.

1.3   Applications and Usage Cases

There are a number of possible applications and usage cases for invertible Bloom lookup tables, some of which we explore here.

Database Reconciliation   Suppose Alice and Bob hold distinct, but similar, copies, D_A and D_B, of an indexed database, D, and they would like to reconcile the differences between D_A and D_B. For example, Alice could hold a current version of D and Bob could hold a backup, or Alice and Bob could represent two different copies of someone's calendar database (say, respectively on a desktop computer and a smartphone) that now need to be symmetrically synchronized. Such usage cases are natural in database systems, particularly for systems that take the approach advocated in an interesting recent CACM article by Stonebraker [36] that argues in favor of sacrificing consistency for the sake of availability and partition-tolerance, and then regaining consistency by performing database reconciliation computations whenever needed. Incidentally, in a separate work, Eppstein *et al.*[2] are currently empirically exploring a similar application of the invertible Bloom table technology for the reconciliation of two distributed mirrors of a filesystem (say, in a peer-to-peer network).

To achieve such a reconciliation with low overhead, Alice constructs an IBLT, $\mathcal{B}$, for D_A, using indices as keys and checksums of her records as values. She then sends the IBLT $\mathcal{B}$ to Bob, who then deletes index-checksum pairs from $\mathcal{B}$ corresponding to all of his entries in D_B. The remaining key-value pairs corresponding to insertions without deletions identify records that Alice has that Bob doesn't have, and the remaining key-value pairs corresponding to deletions without insertions identify records that Bob has that Alice doesn't have. In addition, as we show, Bob can also use $\mathcal{B}$ to identify records that they both possess but with different checksums. In this way, Alice needs only to send a message $\mathcal{B}$ of size $O(t)$, where t here is an upper bound on the number of differences between D_A and D_B, for Bob to determine the identities of their differences (and a symmetric property holds for a similar message from Bob to Alice).

Tracking Network Acknowledgments   As another example application, consider a router $\mathcal{R}$ that would like to track TCP sessions passing through $\mathcal{R}$. In this case, each session corresponds to a key, and may have an associated value, such as the source or the destination, that needs to be tracked. When such flows are initiated in TCP, particular control messages are passed that

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[1]As in the standard RAM model, we assume in this paper that keys and values respectively fit in a single word of memory (which, in practice, could actually be any fixed number of memory words), and we characterize the space used by our data structure in terms of the number of memory words it uses.

[2]Personal communication.

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can be easily detected, allowing the router to add the flow to the structure. Similarly, when a flow terminates, control messages ending the flow are sent. The IBLT supports fast insertions and deletions and can be used to list out the current flows in the system at various times, as long as the number of flows is less than some preset threshold, t. Note this work can be offloaded simply by sending a copy of the IBLT to an offline agent if desired. Furthermore, the IBLT can return the value associated with a flow when queried, with constant probability close to 1. Finally, if at various points, the number of flows spikes above its standard level to well above t, the IBLT will still be able to list out the flows and perform lookups with the appropriate probabilities once the total load returns to t or below. Again, this is a key feature of the IBLT; all keys and values can be reconstructed with high probability whenever the number of keys is below a design threshold, but if the number of keys *temporarily* exceeds this design threshold and later returns to below this threshold, then the functionality will return at that later time.

In this networking setting, sometimes flows do not terminate properly, leaving them in the data structure when they should disappear. Similarly, initialization messages may not be properly handled, leading to a deletion without a corresponding insertion. We show that the IBLT can be modified to handle such errors with minimal loss in performance. Specifically, we can handle keys that are deleted without being inserted, or keys that erroneously obtain multiple values. Even with such errors, we provide conditions for which all valid flows can still all be listed with high probability. Our experimental results also highlight robustness to these types of errors. (Eventually, of course, such problematic keys should be removed from the data structure. We do not concern ourselves with removal policies here; see [6] for some possibilities based on timing structures.)

Oblivious Selection from a Table   As a final motivating application, consider a scenario where Alice has outsourced her data storage needs, including the contents of an important indexed table, $\mathcal{T}$, of size n, to a cloud storage server, Bob, because Alice has very limited storage capacity (e.g., Alice may only have a smartphone). Moreover, because her data is sensitive and she knows Bob is honest-but-curious regarding her data, she encrypts each record of T using a secret key, and random nonces, so that Bob cannot determine the contents of any record from its encryption alone. Such encryptions are not sufficient, however, to fully protect the privacy of Alice's data, as recent attacks show that the way Alice accesses her data can reveal its contents (e.g., see [10]). Alice needs a way of hiding any patterns in the way she accesses her data.

Suppose now that Alice would like to do a simple SELECT query on $\mathcal{T}$ and she is confident that the result will have a size at most t, which is much less than n but still more than she can store locally. Thus, she cannot use techniques from private information retrieval [11, 37], as that would either require storing results back with Bob in a way that could reveal selected indices or using yet another server besides Bob. She could use techniques from recent oblivious RAM simulations [19, 20, 31] to obfuscate her access patterns, but doing so would require $O(n \log^2 n)$ I/Os. Therefore, using existing techniques would be inefficient.

By using an IBLT, on the other hand, she can perfrom her SELECT query much more efficiently. The advantage comes from the fact that an insertion in an IBLT accesses a random set of cells (that is, memory locations) whose addresses depend (via random hash functions) only on the key of the item being inserted. Alice thus uses all the indices for $\mathcal{T}$ as keys, one for each record, and accesses memory as though inserting each record into an IBLT of size $O(t)$. In fact, Alice only inserts those records that satisfy her SELECT query. However, since Alice encrypts each write using a secret key and random nonces, Bob cannot tell when Alice's write operations are actually changing the records stored in the IBLT, and when a write operation is simply rewriting the same contents of a cell over

again re-encrypted with a different nonce. In this way Alice can obliviously create an IBLT of size $O(t)$ that contains the result of her query and is stored by Bob. Then, using existing methods for oblivious RAM simulation [20], she can subsequently obliviously extract the elements from her IBLT using $O(t \log^2 t)$ I/Os. With this approach Bob learns nothing about her data from her access pattern. In addition, the total number of I/Os for her to perform her query is $O(n + t \log^2 t)$, which is linear (and optimal) for any t that is $O(n/\log^2 n)$. We are not currently aware of any other way that Alice can achieve such a result using a structure other than an IBLT.

2   A Simple Version of the Invertible Bloom Lookup Table

In this section, we describe and analyze a simple version of the IBLT. In the sections that follow we describe how to augment and extend this simple structure to achieve various additional performance goals.

The IBLT data structure, $\mathcal{B}$, is a randomized data structure storing a set of key-value pairs. It is designed with respect to a threshold number of keys, t; when we say the structure is successful for an operation with high probability it is under the assumption that the actual number of keys in the structure at that time, which we henceforth denote by n, is less than or equal to t. Note that n can exceed t during the course of normal operation, however.

As mentioned earlier, we assume throughout that, as in the standard RAM model, keys and values respectively fit in a single word of memory (which, in practice, could actually be any fixed number of memory words) and that each such word can alternatively be viewed as an integer, character string, floating-point number, etc. Thus, without loss of generality, we view keys and values as positive integers.

2.1   Operations Supported

Our structure supports the following operations:

- INSERT(x, y): insert the key-value pair, (x, y), into $\mathcal{B}$. This operation always succeeds, assuming that all keys are distinct.
- DELETE(x, y): delete the key-value pair, (x, y), from $\mathcal{B}$. This operation always succeeds, provided $(x, y) \in \mathcal{B}$, which we assume for the rest of this section.
- GET(x): return the value y such that there is a key-value pair, (x, y), in $\mathcal{B}$. If $y =$ null is returned, then $(x, y) \notin \mathcal{B}$ for any value of y. With low (but constant) probability, this operation may fail, returning a "not found" error condition. In this case there may or may not be a key-value pair (x, y) in $\mathcal{B}$.
- LISTENTRIES(): list all the key-value pairs being stored in $\mathcal{B}$. With low (inverse polynomial in t) probability, this operation may return a partial list along with an "list-incomplete" error condition.

When an IBLT $\mathcal{B}$ is first created, it initializes a lookup table T of m cells. Each of the cells in T stores a constant number of fields, each of which corresponds to a single memory word. We emphasize that an important feature of the data structure is that at times the number of key-value pairs in $\mathcal{B}$ can be much larger than m, but the space used for $\mathcal{B}$ remains $O(m)$ words. The INSERT and DELETE methods never fail, whereas the GET and LISTENTRIES methods, on the other hand, only guarantee good probabilistic success when $n \leq t$. For our structures we shall generally have

$m = O(t)$, and often we can give quite tight analyses on the constants required, as we shall see below.

2.2 Data Structure Architecture

Like a standard Bloom filter, an IBLT uses a set of k random[3] hash functions, h_1, h_2, ..., h_k, to determine where items are stored. In our case, each key-value pair, (x, y), is placed into cells $T[h_1(x)]$, $T[h_2(x)]$, ... $T[h_t(x)]$. In what follows, for technical reasons, we assume that the hashes yield distinct locations. This can be accomplished in various ways, with one standard approach being to split the m cells into k subtables each of size m/k, and having each hash function choose one cell (uniformly) from each subtable. Such splitting does not affect the asymptotic behavior in our analysis and can yield other benefits, including ease of parallelization of reads and writes into the hash table. (Another approach would be to select the first k distinct hash values from a specific sequence of random hash functions.)

Each cell contains three fields:

- a count field, which counts the number of entries that have been mapped to this cell,
- a keySum field, which is the sum of all the keys that have been mapped to this cell,
- a valueSum field, which is the sum of all the values that have been mapped to this cell.

Given these fields, which are initially 0, performing the update operations is fairly straightforward:

- INSERT(x, y):

    for each (distinct) $h_i(x)$, for $i = 1, \ldots, k$ **do**
      add 1 to $T[h_i(x)]$.count
      add x to $T[h_i(x)]$.keySum
      add y to $T[h_i(x)]$.valueSum
    end for

- DELETE(x, y):

    for each (distinct) $h_i(x)$, for $i = 1, \ldots, k$ **do**
      subtract 1 from $T[h_i(x)]$.count
      subtract x from $T[h_i(x)]$.keySum
      subtract y from $T[h_i(x)]$.valueSum
    end for

2.3 Data Lookups

We perform the GET operation in a manner similar to how membership queries are done in a standard Bloom filter. The details are as follows:

- GET(x):

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[3]We assume, for the sake of simplicity in our analysis, that the hash functions are fully random, but this does not appear strictly required. For example, the techniques of [24] can be applied if the data has a sufficient amount of entropy. For worst-case data, we are not aware of any results regarding the 2-core of a random hypergraph where the vertices for each edge are chosen according to hash functions with limited independence, which, as we will see, would be needed for such a result. Similar graph problems with limited independence have recently been studied in [1]. It is an interesting theoretical question to obtain better bounds on the randomness needed for our proposed IBLT data structure.

```
for each (distinct) h_i(x), for i = 1, ..., k do
  if T[h_i(x)].count = 0 then
    return null
  else if T[h_i(x)].count = 1 then
    if T[h_i(x)].keySum = x then
      return T[h_i(x)].valueSum
    else
      return null
    end if
  end if
end for
return "not found"
```

Recall that for now we assume that all insertions and deletions are done correctly, that is, no insert will be done for an existing key in $\mathcal{B}$ and no delete will be performed for a key-value pair not already in $\mathcal{B}$. With this assumption, if the above operation returns a value y or the null value, then this is the correct response. Note that this method may fail, returning "not found," if it can find no cell that x maps to that holds only one entry.

For a key x that is in $\mathcal{B}$, consider the probability p_0 that each of its hash locations contains no other item. Using the standard analysis for Bloom filters (e.g., see [8]), we find p_0 is:

$$p_0 = \left(1 - \frac{k}{m}\right)^{(n-1)} \approx e^{-kn/m}.$$

That is, assuming the table is split into k subtables of size m/k (one for each hash function), each of the other $n-1$ keys misses the location independently with probability $1 - k/m$. One nice interpretation of this is that the number of keys that hash to the cell is approximately a Poisson random variable with mean kn/m, and $e^{-kn/m}$ is the corresponding probability a cell is empty. The probability that a GET for a key that is in $\mathcal{B}$ returns "not found" is therefore approximately

$$(1 - p_0)^k \approx \left(1 - e^{-kn/m}\right)^k,$$

which corresponds to the false-positive rate for a standard Bloom filter. As is standard for these arguments, these approximations can be readily replaced by tight concentration results [8].

The probability that a GET for a key that is *not* in $\mathcal{B}$ returns "not found" instead of null can be found similarly. Here, however, note that every cell hashed to by that key must be hashed to by at least two other keys from $\mathcal{B}$. This is because an empty cell would lead to a null return value, and a cell with just one key hashed to it would yield the corresponding true key value, and hence also lead to a null return value for a key not in $\mathcal{B}$. Using the same approximation – specifically, that the number of keys from $\mathcal{B}$ that land in a cell is approximately distributed as a discrete Poisson random variable with mean kn/m – we find this probability is

$$\left(1 - e^{-kn/m} - \frac{kn}{m}e^{-kn/m}\right)^k.$$

2.4   Listing Set Entries

Let us next consider the method for listing the contents of $\mathcal{B}$. We describe this method in a destructive fashion—if one wants a non-destructive method, then one should first create a copy of $\mathcal{B}$ as a backup.

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- LISTENTRIES():
    - **while** there's an $i \in [1, m]$ with $T[i]$.count $= 1$ **do**
        - add the pair $(T[i]$.keySum$, T[i]$.valueSum$)$ to the output list
        - call DELETE$(T[i]$.keySum$, T[i]$.valueSum$)$
    - **end while**

It is a fairly straightforward exercise to implement this method in $O(m)$ time, say, by using a link-list-based priority queue of cells in T indexed by their count fields and modifying the DELETE method to update this queue each time it deletes an entry from $\mathcal{B}$.

If at the end of the while-loop all the entries in T are empty, then we say that the method *succeeded* and we can confirm that the output list is the entire set of entries in $\mathcal{B}$. If, on the other hand, there are some cells in T with non-zero counts, then the method only outputs a partial list of the key-value pairs in $\mathcal{B}$.

This process should appear entirely familiar to those who work with random graphs and hypergraphs. It is exactly the same procedure used to find the 2-core of a random hypergraph (e.g., see [16, 25]). To make the connection, think of the cells as being vertices in the hypergraph, and the elements as being hyperedges, with the vertices for an edge corresponding to the hash locations for the element. The 2-core is the largest sub-hypergraph that has minimum degree at least 2. The standard "peeling process" finds the 2-core: while there exists a vertex with degree 1, delete it and the corresponding hyperedge. The equivalence between the peeling process and the scheme for LISTENTRIES is immediate. We note that this peeling process is similarly used for various erasure-correcting codes, such as Tornado codes and its derivatives (e.g., see [23]), that have, in some ways, the same flavor as this construction[4].

Assuming that the cells associated with an element are chosen uniformly at random, we use known results on 2-cores of random hypergraphs. In particular, tight thresholds are known; when the degree k of each element is at least 2, there are constants $c_k > 1$ such that if $m > (c_k + \epsilon)n$ for any constant $\epsilon > 0$, LISTENTRIES succeeds with high probability, that is with probability $1 - o(1)$. Similarly, if $m < (c_k - \epsilon)n$ for any constant $\epsilon > 0$, LISTENTRIES succeeds with probability $o(1)$. As can be found in for example [16, 25], these values are given by

$$c_k^{-1} = \sup \left\{ \alpha : 0 < \alpha < 1; \forall x \in (0, 1), 1 - e^{-k\alpha x^{k-1}} < x \right\}.$$

The numerical values for these thresholds for $3 \leq k \leq 7$ are given in Table 1, although here we are not truly concerned with the exact values c_k; it is enough that only linear space is required. It is worthwhile noting that c_k is generally close to 1, while to obtain successful GET operations we require a number of cells which is a significant constant factor larger than n. Therefore, in practice the choice of the size of the IBLT will generally be determined by the desired probability for a successful GET operation, not the need for listing.

For our desired applications, we want to be clear about the high probability of failure when listing entries; specifically, we may desire it to be $O(n^{-c})$ for a chosen constant c. By choosing k sufficiently large and m above the 2-core threshold, we can ensure this; indeed, standard results give that the bottleneck is the possibility of having two edges with the same collection of vertices,

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[4]Following this analogy, one could for example, consider *irregular* versions of the IBLT, where different keys utilize a different number of hash values; such a variation could use less space while allowing LISTENTRIES to succeed, or could be used to allow some keys with more hash locations to obtain a better likelihood of a successful lookup. These variations are straightforward and we do not consider the details further here.

k	3	4	5	6	7
c_k	1.222	1.295	1.425	1.570	1.721

Table 1: Thresholds for the 2-core rounded to four decimal places.

giving a failure probability of $O(n^{-k+2})$. The following theorem follows from previous work but we provide it for completeness.

Theorem 1: *As long as m is chosen so that $m > (c_k + \epsilon)n$ for some $\epsilon > 0$, LISTENTRIES fails with probability $O(n^{-k+2})$.*

Proof: We describe the result in terms of the 2-core. The probability that j hyperedges form a non-empty 2-core is dominated by the probability that these edges utilize only $jk/2$ vertices. This probability is at most

$$\binom{n}{j}\binom{m}{jk/2}\left(\frac{jk}{2m}\right)^{jk} \leq \left(\frac{n}{e}\right)^j \left(\frac{2m}{jk}\right)^{jk/2}\left(\frac{jk}{2m}\right)^{jk}$$

$$= \frac{n^j}{m^{jk/2}}\left(\frac{jke}{m}\right)^{jk/2}\left(\frac{e}{j}\right)^j.$$

For k constant, $m > (c_k + \epsilon)n$, and $j \leq \gamma n$ for some constant γ, the sum of these probabilities is dominated by the term where $j = 2$, which corresponds to a failure probability of $O(n^{-k+2})$. To deal separately with the case of $j > \gamma n$, we note that standard analysis of the peeling process shows that, as long as m is above the decoding threshold, the probability that the peeling process fails before reaching a core of size δn for any constant δ is asymptotically exponentially small in n. (See, e.g., [14].) By this argument the case of $j > \gamma n$ adds a vanishing amount to the failure probability, completing the proof of the theorem. ∎

3 Adding Fault Tolerance to an Invertible Bloom Lookup Table

For cases where there can be deletions for key-value pairs that are not already in $\mathcal{B}$, or values can be inserted for keys that are already in $\mathcal{B}$, we require some fault tolerance. We can utilize a standard approach of adding random checksums to get better fault tolerance.

Extraneous Deletions Let us first consider a case with extraneous deletions only. Specifically, we assume a key-value pair might be deleted without a corresponding insertion; however, in this first setting we still assume each key is associated with a single value, and is not inserted or deleted multiple times at any instant. This causes a variety of problems for both the GET and LISTENTRIES routines. For example, it is possible for a cell to have an associated count of 1 even if more than one key has hashed to it, if there are corresponding extraneous deletions; this causes us to re-evaluate our LISTENTRIES routine.

To help deal with these issues, we add to our IBLT structure. We assume that each key x has an additional hash value given by a hash function $G_1(x)$, which in general we assume will take on uniform random values in a range $[1, R]$. We then require each cell has the following additional field:

- a hashkeySum field, which is the sum of the hash values, $G_1(x)$, for all the keys that have been mapped to this cell.

The hashkeySum field must be of sufficiently many bits and the hash function must be sufficiently random to make collisions sufficiently unlikely; this is not hard to achieve in practice. Our insertion and deletion operations must now change accordingly, in that we now must add $G_1(x)$ to each $T[h_i(x)]$.hashkeySum on an insertion and subtract $G_1(x)$ during a deletion. The pseudocode for these and the other operations is given in Figure 3 at the end of this paper.

The hashkeySum field can serve as an extra check. For example, to check when a cell has a count of 1 that it corresponds to a cell without extraneous deletions, we check $G_1(x)$ field against the hashkeySum field. For an error to occur, we must have that a deletion has caused a count of 1 where the count should be higher, and the hashed key values must align so that their sum causes a false check. This probability is clearly at most $1/R$ (using the standard principle of deferred decisions, the "last hash" must take on the precise wrong value for a false check). We will generally assume that R is chosen large enough that we can assume a false match does not occur throughout the lifetime of the data structure, noting that only $O(\log n)$ bits are needed to handle lifetimes that are polynomial in n.

Let us now consider GET operations. The natural approach is to assume that the hashkeySum field will not lead to a false check, as above. In this case, on a GET of a key x, if the count field is 0, and the keySum and hashkeySum are also 0, one should assume that the cell is in fact empty, and return null. Similarly, if the count field is 1, and the keySum and hashkeySum match x and $G_1(x)$, respectively, then one should assume the cell has the right key, and return its value. In fact, if the count field is -1, and after negating keySum and hashkeySum the values match x and $G_1(x)$, respectively, one should assume the cell has the right key, except that it has been deleted instead of inserted! In our pseudocode we return the value, although one could also flag it as an extraneous deletion as well. Note, however, that we can no longer return null if the count field is 1 but the keySum field does not match x; in this case, there could be, for example, an additional key inserted and an additional key extraneously deleted from that cell, which would cause the field to not match even if x was hashed to that cell. If we let n be the number of keys either inserted or extraneously deleted in the IBLT, then this reduces the probability of returning null for a key not in $\mathcal{B}$ to $\left(1 - e^{-kn/m}\right)^k$. That is, to return null we must have at least one cell with zero items from $\mathcal{B}$ hashing to it, which occurs (approximately) with the given probability (using our Poisson approximation).

For the LISTENTRIES operation, we again use hashkeySum field to check when a cell has a count of 1 that it corresponds to a cell without extraneous deletions. Note here that an error in this check will cause the entire listing operation to fail, so the probability of a false check should be made quite low—certainly inverse polynomial in n. Also note, again, that we can make progress in recovering keys with cells with a count of -1 as well, if the cell contains only one extraneously deleted key and no inserted keys. That is, if a cell contains a count of -1, we can negate the count, keySum, and hashkeySum fields, check the hash value against the key to prevent a false match, and if that check passes recover the key and remove it (in this case, add it back in) to the other associated cells. Hence, a cell cannot yield a key during the listing process only if more than one key, either inserted or deleted, has hashed to that cell. This is now exactly the same setting as in the original case of no extraneous deletions, and hence (assuming that no false checks occur!) the same analysis applies, with n representing the number of keys either inserted or extraneously deleted. We give the revised pseudo-code descriptions in Figure 3.

Multiple Values A more challenging case for fault tolerance occurs when a key can be inserted multiple times with different values, or inserted and deleted with different values. If a key is

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inserted multiple times with different values, not only can that key not be recovered, but every cell associated with that key has been *poisoned*, in that it will not be useful for listing keys, as it cannot have a count of 1 even as other values are recovered. (A later deletion of a key-value pair could correct this problem, of course, but the cell is poisoned at the time.) The same is true if a key is inserted and deleted with different values, and here the problem is potentially even worse: if a single other key hashes to that cell, the count may be 1 and the keySum and hashkeySum fields will be correct even though the valueSum field will not match the other key's value, causing errors.

Correspondingly, we introduce an additional check for the sum of the values at a cell, using a hash function $G_2(y)$ for the values, and adding the following field:

- a hashvalueSum field, which is the sum of the hash values $G_2(y)$ for all the values that have been mapped to this cell.

One can then check that the hash of the keySum and valueSum take on the appropriate values when the count field of a cell is 1 (or -1) in order to see if listing the key-value pair is appropriate.

The question remains whether the poisoned cells will prevent recovery of key values. Here we modify the goal of LISTENTRIES to return all key-value pairs for all valid keys with high probability—that is, all keys with a single associated value at that time. We first claim that if the invalid keys make up a constant fraction of the n keys that this is not possible under our construction with linear space. A constant fraction of the cells would then be poisoned, and with constant probability each valid key would then hash solely to poisoned cells, in which case the key could not be recovered.

However, it is useful to consider these probabilities, as in practical settings these quantities will determine the probability of failure. For example, suppose γn keys are invalid for some constant γ. By our previous analysis, the fraction of cells that are poisoned is concentrated around $\left(1 - e^{-k\gamma n/m}\right)$, and hence the probability that any specific valid key has all of its cells poisoned is $\left(1 - e^{-k\gamma n/m}\right)^k$. (While there are other possible ways a key could not be recovered, for example if two keys have all but one of their cells poisoned and their remaining cell is the same, this gives a good first approximation for reasonable values, as other terms will generally be lower order when these probabilities are small.) For example, in a configuration we use in our experiments below, we choose $k = 5$, $m/n = 8$, and $\gamma = 1/10$; in this case, the probability of a specific valid key being unrecoverable is approximately $8.16 \cdot 10^{-7}$, which may be quite suitable for practice.

For a more theoretical asymptotic analysis, suppose instead that there are only $n^{1-\beta}$ invalid keys. Then if each key uses at least $\lceil 1/\beta \rceil + 1$ hash functions, with high probability every valid key will have at least one hash location that does not coincide with a invalid cell. This alone does not guarantee reconstructing the valid key-value pairs, but we can extend this idea to show LISTENTRIES can provably successfully obtain all valid keys even with $n^{1-\beta}$ invalid keys; by using $k = \lceil 1/\beta \rceil + 4$ hash functions, we can guarantee with high probability that every valid key has at least 3 hash locations without an invalid cell. (One can raise the probability to any inverse polynomial in n by changing the constant 4 as desired.) Indeed, we can determine the induced distribution for the number of neighbors to unpoisoned cells for the valid keys, but the important fact is that the number of keys with with k hashes to unpoisoned cells in $n - o(n)$. It follows from the standard analysis of random cores (e.g., see Molloy [25]) that the same threshold as for the original setting with k hash functions applies. Hence the number of cells needed will again be linear in n (with the coefficient dependent on β) in order to guarantee successful listing of keys with high probability. This yields the following theorem:

Theorem 2: *Suppose there are $n^{1-\beta}$ invalid keys. Let $k = \lceil 1/\beta \rceil + 4$. Then if $m > (c_k + \epsilon)n$ for some $\epsilon > 0$,* LISTENTRIES *succeeds with high probability.*

While this asymptotic analysis provides some useful insights, namely that full recovery is practical, in practice we expect the analysis above based on setting γ so that the number of invalid keys is γn will prove more useful for parameter setting and predicting performance.

Extensions to Duplicates   Interestingly, using the same approach as for extraneous deletions, our IBLT can handle the setting where a key-value pair is inserted multiple times with the *same* value for the key each time. Essentially, this means the IBLT is robust to duplicates, or can also be used to obtain a count for key-value pairs that are inserted multiple times. We again use the additional hashkeySum and valueSum fields. When the count field is j for the cell, we take the keySum, hashkeySum, and valueSum fields and divide them by j to obtain the proposed key, value, and corresponding hash. If the key hash matches, we assume that we have found the right key and return the lookup value or list the key-value pair accordingly, depending on whether a GET or LISTENTRIES operation is being performed. If it is possible to have the same key appear with multiple values, as above, then we must *also* make use of the hashvalueSum fields, dividing it by j and using it to check that the value is correct as well. For the listing operation, the IBLT deletes j copies of the key-value pair from the other cells.[5] The point here is that a key that appears multiple times, just as a key that is deleted rather than inserted, can be handled without significant change to the decoding approach.

Fault Tolerance to Lost Memory Subblocks   We offer one additional way this structure proves resilient to various possible faults. Suppose that the structure is indeed set up with k different memory subblocks, one for each hash function. Conceivably, one could lose an entire subblock, and still be able to recover all the keys in a listing with high probability, with only a reduction in the success probability of a GET (as long as $k - 1$ hashes with a smaller range space remains sufficient for listing). In some sense, this is because the system is arguably overdesigned; obtaining high lookup success probability when n is less than the threshold t requires a large number of empty cells, and this space is far more than is needed for decoding.

An Example Application   As an example of where these ideas might be used, we return to our mirror site application. An IBLT $\mathcal{B}$ from Alice can be used by Bob to find filename-checksum (key-value) pairs where his filename has a different checksum than Alice's. After deleting all his key-value pairs, he lists out the contents of $\mathcal{B}$ to find files that he or Alice has that the other does not. The IBLT might not be empty at this point, however, as the listing process might not have been able to complete due to *poisoned* cells, where deletions were done for keys with values different than Alice's values. To discover these, Bob can re-insert each of his key-value pairs, in turn, to find any that may unpoison a cell in $\mathcal{B}$ (where he immediately deletes ones that don't lead to a new unpoisoned cell). If a new unpoisoned cell is found found (using the G_1 hash function as a check), then Bob can then remove a key-value pair with the same key as his but with a different value (that is, with Alice's value). Note Bob may then also be able to possibly perform more listings of keys that might have been previously unrecovered because of the poisoned cells. Repeating this will discover with high probability all the key-value pairs where Alice and Bob differ.

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[5]Note that here we are making use of the assumption that the hash locations are distinct for a key; otherwise, the count for the number of copies at this location might not match the number of copies of the key in all the other locations.

4 Space-Saving for an Invertible Bloom Lookup Table

Up to this point, we have not been concerned with minimizing space for our IBLT structure, noting that it can be done in linear space. Nevertheless, there are a variety of techniques available for reducing the space required, generally at the expense of additional computation and shuffling of memory, while still keeping constant amortized or worst-case time bounds on various operations. Whether such efforts are worthwhile depends on the setting; in some applications space may be the overriding issue, while in others speed or even simplicity might be more relevant. We briefly point to some of the previous work that can offer further insights.

IBLTs, like other Bloom filter structures, often have a great deal of wasted space corresponding to zero entries that can be compressed or fixed-length space required for fields like the count field that can be made variable-length depending on the value. This wasted space can be compressed away using techniques for keeping compressed forms of arrays, including those for storing arrays of variable-length strings. Such mechanisms are explored in for example [4, 28, 34] and can be applied here.

A simpler, standard approach (saving less space) is to use quotienting, whereby the hash value for a key determines a bucket and an additional quotient value that can be stored. Quotienting can naturally be used with the IBLT to reduce the space used for storing for example the keySum or the hashkeySum values. Another simplification when in a setting where duplicate keys will not appear is to use an XOR in place of a sum operation to store sums of keys and values (or their hashes); as long as the same key does not appear more than once the algorithms presented will continue to apply.

Finally, we recall that the space requirements arise because of the desire for high accuracy for GET operations, not because of the LISTENTRIES operation. If one is willing to give up lookup accuracy—which may be the case if, for example, one expects the system to be overloaded much of the time—then less space is needed to maintain successful listing.

5 Simulations and Experiments

We have run a number of simulations to test the IBLT structure and our analysis. In these experiments we have not focused on running time; a practical implementation could require significant optimization. Instead our focus in on the success probability of the listing of keys and, to a lesser extent, on the success probability for a GET operation. Overall, we have found that the IBLT works quite effectively and the performance matches our theoretical analysis. We provide a few example results. In all of the experiments here, we have chosen to use five hash functions.

First, our calculated asymptotic thresholds for decoding from Table 1 are quite accurate even for reasonably small values. For example, in the setting where there are no duplicate keys or extraneous deletions of keys, we repeatedly performed 20,000 simulations with 10,000 items, and varied the number of cells. Table 1 suggests an asymptotic threshold for listing all entries near 14,250. As shown in Figure 1(a), around this point we see a dramatic increase in the average number of items recovered when performing our LISTENTRIES operation. In fact, at 14,500 cells only two trials failed to recover all key-value pairs, and with 14,600 cells or more all trials successfully recover all key-value pairs. We performed an additional 200,000 trials with 14,600 cells, and again all trials succeeded. In Figure 1(b), we consider 20,000 simulations with 100,000 items, where the corresponding threshold should be near 142,500. With more items, we expect tighter concentration around the threshold, and indeed with 144,000 cells or more all trials successfully recover all key-
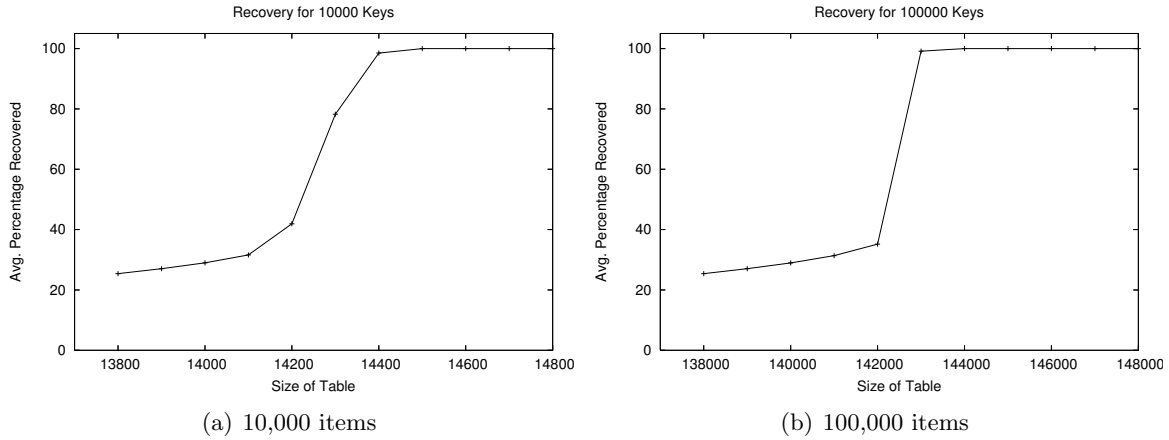
Figure 1: Percentage of items recovered around the threshold. Slightly over the theoretical asymptotic threshold, we obtain full recovery of all items with LISTENTRIES on all simulations. Each data point represents the average of 20,000 simulations.

value pairs. We performed an additional 200,000 trials with with 144,000 cells, and again all trials succeeded.

We acknowledge that more simulations would be required to obtain detailed bounds on the probability of failure to recover all key-value pairs for specific values of the number of items and cells. This is equivalent to the well-studied problem of "finite-length analysis" for related families of error-correcting codes. Dynamic programming techniques, as discussed in [21] and subsequent follow-on work, can be applied to obtain such bounds.

Our next tests of the IBLT allow duplicate keys with the same value and extraneous deletions, but without keys with multiple values. Our analysis suggests this should work exactly as with no duplicate or extraneous deletions, and our simulations verify this. In these simulations, we had each key result in a duplicate with probability 1/5, and each key result in a deletion in place of an insertion with probability 1/5. Using a check on key and value fields, in 20,000 simulations with 10,000 keys, 80,000 cells, and 5 hash functions, a complete listing was obtained every time, and GET operations were successful on average for 97.83 percent of the items, matching the standard analysis for a Bloom filter. Results were similar with 20,000 runs with 100,000 keys and 800,000 cells, again with complete recovery each time and get operations successful on average for 97.83 percent of the keys.

Finally, we tested the IBLT with keys that erroneously obtain multiple values. As expected, these keys can both prevent recovery of other items during listing, but do not impact the success probability of GET operations for other items. For example, again using a check on key and value fields, in 20,000 simulations with 10,000 keys of which 500 had multiple values, 80,000 cells, and 5 hash functions, the 9500 remaining key-value pairs were recovered 19,996 times; the remaining 4 times all but one of the 9500 items was recovered. With 1000 keys with multiple values, the remaining key-value pairs were recovered 19,872 times, and again the remaining 128 times all but one of the 9000 items was recovered. The average success rate for GET operations remained 97.83 percent on the valid keys in both cases, as would be expected. We note that 10,000 keys and 1,000 keys with multiple values, our previous back-of-the-envelope calculation showed that each valid key
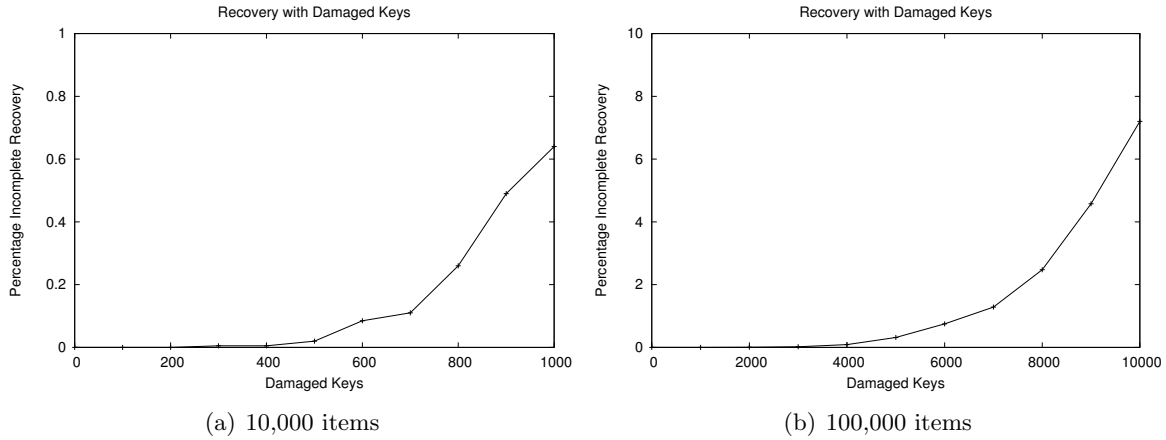
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| (a) 10,000 items | (b) 100,000 items |

Figure 2: Percentage of trials with incomplete recovery with "damaged" keys that have multiple values. Each data point represents the average of 20,000 simulations.

would fail with probability roughly $8.16 \cdot 10^{-7}$; hence, with 9,000 other keys, assuming independence we would estimate the probability of complete recovery at approximately 0.9927, closely matching our experimental results. More detailed results are give in Figures 2(a) and 2(b), where we vary the number keys with multiple values for two settings: 10,000 keys and 80,000 cells, and 100,000 keys and 800,000 cells. The results are based on 20,000 trials. As can be seen complete recovery is possible with large numbers of multiple-valued keys in both cases, but naturally the probability of complete recovery becomes worse with larger numbers of keys even if the percentage of invalid keys is the same.

We emphasize that even when complete recovery does not occur in this setting, generally almost all keys with a single value can be recovered. For example, in Table 2 we consider three experiments. The first is for 10,000 keys, 80,000 cells, and 1,000 keys with duplicate values. The second is the same but with 2,000 keys with duplicate values. The third is for 100,000 keys, 800,000 cells, and 10,000 keys with duplicate values. Overall 20,000 trials for each experiment, in no case were more than 3 valid keys unrecovered. The main point of Table 2 is that with suitable design parameters, even when complete recovery is not possible because of invalid keys, the degradation is minor. We suspect this level of robustness may be useful for some applications where almost-complete recovery is acceptable.

	Unrecovered Keys			
	0	1	2	3
Experiment 1	99.360	0.640	0.000	0.000
Experiment 2	83.505	14.885	1.520	0.090
Experiment 3	92.800	6.915	0.265	0.020

Table 2: Percentage of trials where 1, 2, and 3 keys are left unrecovered.

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6   Conclusion and Future Work

We have given an extension to the Bloom filter data structure to key-value pairs and the ability to list out its contents. This structure is deceptively simple, but is able to achieve functionalities and efficiencies that appear to be unique in many respects, based on our analysis derived from recent results on 2-cores in hypergraphs. One possible direction for future work includes whether one can easily include methods for allowing for multiple values as a natural condition instead of an error.

Acknowledgments

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- INSERT(x, y):

  for each $h_i(x)$ value, for $i = 1, \ldots, k$ **do**

      add 1 to $T[h_i(x)]$.count

      add x to $T[h_i(x)]$.keySum

      add y to $T[h_i(x)]$.valueSum

      add $G_1(x)$ to $T[h_i(x)]$.hashkeySum

  end for

- DELETE(x, y):

  for each $h_i(x)$ value, for $i = 1, \ldots, k$ **do**

      subtract 1 from $T[h_i(x)]$.count

      subtract x from $T[h_i(x)]$.keySum

      subtract y from $T[h_i(x)]$.valueSum

      subtract $G_1(x)$ to $T[h_i(x)]$.hashkeySum

  end for

- GET(x):

  for each $h_i(x)$ value, for $i = 1, \ldots, k$ **do**

      if $T[h_i(x)]$.count $= 0$ **and** $T[h_i(x)]$.keySum $= 0$ **and** $T[h_i(x)]$.hashkeySum $= 0$ **then**

        return null

      else if $T[h_i(x)]$.count $= 1$ **and** $T[h_i(x)]$.keySum $= x$ **and** $T[h_i(x)]$.hashkeySum $= G_1(x)$
  then

        return $T[h_i(x)]$.valueSum

      else if $T[h_i(x)]$.count $= -1$ **and** $T[h_i(x)]$.keySum $= -x$ **and** $T[h_i(x)]$.hashkeySum $=
  -G_1(x)$ **then**

        return $-T[h_i(x)]$.valueSum

      end if

  end for

  return "not found"

- LISTENTRIES():

  while there is an $i \in [1, m]$ such that $T[i]$.count $= 1$ or $T[i]$.count $= -1$ **do**

      if $T[h_i(x)]$.count $= 1$ **and** $T[h_i(x)]$.hashkeySum $= G_1(T[h_i(x)]$.keySum) **then**

        add the pair, $(T[i]$.keySum$, T[i]$.valueSum$)$, to the output list

        call DELETE$(T[i]$.keySum$, T[i]$.valueSum$)$

      else if $T[h_i(x)]$.count $= -1$ **and** $-T[h_i(x)]$.hashkeySum $= G_1(-T[h_i(x)]$.keySum$)$
  then

        add the pair, $(-T[i]$.keySum$, -T[i]$.valueSum$)$, to the output list

        call INSERT$(-T[i]$.keySum$, -T[i]$.valueSum$)$

      end if

  end while

Figure 3: Revised pseudo-code for tolerating extraneous deletions.