Cache-Oblivious Peeling of Random Hypergraphs

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Abstract

The computation of a peeling order in a randomly generated hypergraph is the most time-consuming step in a number of constructions, such as perfect hashing schemes, random r-SAT solvers, error-correcting codes, and approximate set encodings. While there exists a straightforward linear-time algorithm, its poor I/O performance makes it impractical for hypergraphs whose size exceeds the available internal memory.

We show how to reduce the computation of a peeling order to a small number of sequential scans and sorts, and analyze its I/O complexity in the cache-oblivious model. The resulting algorithm requires $O(\mathrm{sort}(n))$ I/Os and $O(n \log n)$ time to peel a random hypergraph with n edges.

We experimentally evaluate the performance of our implementation of this algorithm in a real-world scenario by using the construction of minimal perfect hash functions (MPHF) as our test case: our algorithm builds a MPHF of 7.6 billion keys in less than 21 hours on a single machine. The resulting data structure is both more space-efficient and faster than that obtained with the current state-of-the-art MPHF construction for large-scale key sets.

1 Introduction

Hypergraphs can be used to model sets of dependencies among variables of a system: vertices correspond to variables and edges to relations of dependency among variables, such as equations binding variables together. This correspondence can be used to transfer graph-theoretical properties to solvability conditions in the original system of dependencies.

Among these, one of the most useful is the concept of peeling order. Given an r-hypergraph, a *peeling order* is an order of its edges such that each edge has a vertex of degree 1 in the subgraph obtained by removing the previous edges in the order. Such an order exists if the hypergraph does not have a non-empty 2-core, that is, a set of vertices that induces a subgraph whose vertices have all degree at least 2.

In the above interpretation, if the equations of a system are arranged in peeling order, then each equation has at least one variable that does not appear in any equation that

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comes later in the ordering, i.e., the system becomes *triangular*, so it can be easily solved by backward substitution. For this reason, peeling orders have found applications in a number of fundamental problems, such as hash constructions [3, 6, 8, 9, 16], solving random instances of r-SAT [9, 19, 20], and the construction of error-correcting codes [11, 15, 18]. These applications exploit the guarantee that if the edge sparsity γ of a random r-hypergraph is larger than a certain sparsity threshold c_r (e.g., $c_3 \approx 1.221$), then w.h.p. (with high probability) the hypergraph has an empty 2-core [20].

The construction of *perfect hash functions* (PHF) is probably the most important of the aforementioned applications. Given a set S of n keys, a PHF for S maps injectively the keys into the set of the first $m \geq n$ natural numbers. A perfect hash function is *minimal* (MPHF) if $m = n$. A lower bound by Mehlhorn [17] states that $n \log e \approx 1.44n$ bits are necessary to represent a MPHF; a matching (up to lower order terms) upper bound is provided in [12], but the construction is impractical. Most practical approaches, instead, are based on random 3-hypergraphs, resulting in MPHFs that use about $2c_3n \approx 2.5n$ bits [6, 8, 16]. These solutions, which we review in Section 3, build on the MWHC technique [16], whose most demanding task is in fact the computation of a peeling order.

There is a surprisingly simple greedy algorithm to find a peeling order when it exists, or a 2-core when it does not: find a vertex of degree 1, remove (*peel*) its only edge from the hypergraph, and iterate this process until either no edges are left (in which case the removal order is a peeling order), or all the non-isolated vertices left have degree at least 2 (thus forming a 2-core). This algorithm can be easily implemented to run in linear time and space.

MPHFs are the main ingredient in many space-efficient data structures, such as (compressed) full-text indexes [4], monotone MPHFs [1], Bloom filter-like data structures [5], and prefix-search data structures [2].

It should be clear that the applications that benefit the most from such data structures are those involving large-scale key sets, often orders of magnitude larger than the main memory. Unfortunately, the standard linear-time peeling algorithm requires several tens of bytes per key of working memory, even if the final data structure can be stored in just a handful of bits per key. It is hence common that, while the data structure fits in memory, such memory is not enough to actually *build* it. It is then necessary to resort to external memory, but the poor I/O performance of the algorithm makes such an approach impossible.

Application-specific workarounds have been devised; for example, Botelho et al. [6] proposed an algorithm (called HEM) to build MPHFs in external memory by splitting the key set into small buckets and computing independent MPHFs for each bucket. A first-level index is used to find the bucket of a given key. The main drawback of this solution is that the first-level index introduces a non-negligible overhead in both space and lookup time; moreover, this construction cannot be extended to applications other than hashing.

In this paper we provide the first efficient algorithm in the *cache-oblivious* model that, given a random r-hypergraph with n edges and γn vertices (with $r = O(1)$ and $\gamma > c_r$), computes a peeling order in time $O(n \log n)$ and with $O(\mathrm{sort}(n))$ I/Os w.h.p., where $\mathrm{sort}(n)$ is the I/O complexity of sorting n keys. By applying this result we can construct (monotone) MPHFs, static functions, and Bloom filter-like data structures in $O(\mathrm{sort}(n))$ I/Os. In our experimental evaluation, we show that the algorithm makes it indeed possible to peel very large hypergraphs: an MPHF for a set of 7.6 billion keys is computed in less than 21 hours; on the same hardware, the standard algorithm would not be able to manage more than 2.1

billion keys. Although we use minimal perfect hash functions construction as our test case, results of these experiments remain valid for all the other applications due to the random nature of the underlying hypergraphs.

2 Notation and tools

Model and assumptions We analyze our algorithms in the cache-oblivious model [10]. In this model, the machine has a two-level memory hierarchy, where the fast level has an unknown size of M words and a slow level of unbounded size where our data reside. We assume that the fast level plays the role of a cache for the slow level with an optimal replacement strategy where the transfers (a.k.a. I/Os) between the two levels are done in blocks of an unknown size of $B \leq M$ words; the I/O cost of an algorithm is the total number of such block transfers. *Scanning* and *sorting* are two fundamental building blocks in the design of cache-oblivious algorithms [10]: under the tall-cache assumption [7], given an array of N contiguous items the I/Os required for scanning and sorting are $\mathrm{scan}(N) = O(1 + N/B)$ and $\mathrm{sort}(N) = O(N/B \log_{M/B}(N/B))$ I/Os.

Hypergraphs An r-hypergraph on a vertex set V is a subset E of $\binom{V}{r}$, the set of subsets of V of cardinality r. An element of E is called an *edge*. We call an ordered r-tuple from V an *oriented edge*; if e is an edge, an oriented edge whose vertices are those in e is called an *orientation* of e. From now on we will focus on 3-hypergraphs; generalization to an arbitrary integer r is straightforward. We define *valid* orientations those oriented edges (v_0, v_1, v_2) where $v_1 < v_2$ (for arbitrary r, $v_1 < \cdots < v_{r-1}$). Then for each edge there are 6 orientations, but only 3 valid orientations ($r!$ orientations of which r are valid).

We say that a valid oriented edge (v_0, v_1, v_2) is the i-th orientation if v_0 is the i-th smallest among the three; in particular, the 0-th orientation is the *canonical* orientation. Edges correspond bijectively with their canonical orientations. Furthermore, valid orientations can be mapped bijectively to pairs (e, v) where e is an edge and v a vertex contained in e, simply by the correspondence $(v_0, v_1, v_2) \mapsto (\{v_0, v_1, v_2\}, v_0)$. In the following all the orientations are assumed to be valid, so we will use the term *orientation* to mean *valid orientation*.

3 The Majewski–Wormald–Havas–Czech technique

Majewski et al. [16] proposed a technique (MWHC) to compute an *order-preserving minimal perfect hash function*, that is, a function mapping a set of keys S in some specified way into $[|S|]$. The technique actually makes it possible to store succinctly any function $f : S \to [\sigma]$, for arbitrary σ. In this section we briefly describe their construction.

First, we choose three random[1] hash functions $h_0, h_1, h_2 : S \to [\gamma n]$ and generate a 3-hypergraph[2] with γn vertices, where γ is a constant above the *critical threshold* c_3, by mapping each key x to the edge $\{h_0(x), h_1(x), h_2(x)\}$. The goal is to find an array u of γn integers in $[\sigma]$ such that for each key x one has $f(x) = (u_{h_0(x)} + u_{h_1(x)} + u_{h_2(x)}) \mod \sigma$. This yields a linear system with n equations and γn variables u_i; if the associated hypergraph

[1] Like most MWHC implementations, in our experiments we use a Jenkins hash function with a 64-bit seed in place of a fully random hash function.

[2] Although the technique works for r-hypergraphs, $r = 3$ provides the lowest space usage [20].

is peelable, it is easy to solve the system. Since γ is larger than the critical threshold, the algorithm succeeds with probability $1 - o(1)$ as $n \to \infty$ [20].

By storing such values u_i, each requiring $\lceil \log \sigma \rceil$ bits, plus the three hash functions, we will be able to recover $f(x)$. Overall, the space required will be $\lceil \log \sigma \rceil \gamma n$ bits, which can be reduced to $\lceil \log \sigma \rceil n + \gamma n + o(n)$ using a ranking structure [13]. This technique can be easily extended to construct MPHFs: we define the function $f \colon S \to [3]$ as $x \mapsto i$ where $h_i(x)$ is a degree-1 vertex when the edge corresponding to x is peeled; it is then easy to see that $h_{f(x)}(x) : S \to [\gamma n]$ is a PHF. The function can be again made minimal by adding a ranking structure on the vector u [6].

As noted in the introduction, the peeling procedure needed to solve the linear system can be performed in linear time using a greedy algorithm (referred to as *standard linear-time peeling*). However, this procedure requires random access to several integers per key, needed for bookkeeping; moreover, since the graph is random, the visit order is close to random. As a consequence, if the key set is so large that it is necessary to spill to the disk part of the working data structures, the I/O volume slows down the algorithm to unacceptable rates.

Practical workarounds (HEM) Botelho et al. [6] proposed a practical external-memory solution: they replace each key with a *signature* of $\Theta(\log n)$ bits computed with a random hash function, so that no collision occurs. The signatures are then sorted and divided into small buckets based on their most significant bits, and a separate MPHF is computed for each bucket with the approach described above. The representations of the bucket functions are then concatenated into a single array and their offsets stored in a separate vector.

The construction algorithm only requires to sort the signatures (which can be done efficiently in external memory) and to scan the resulting array to compute the bucket functions; hence, it is extremely scalable. The extra indirection needed to address the blocks causes however the resulting data structure to be both *slower* and *larger* than one obtained by computing a single function on the whole key set. In their experiments with a practical version of the construction, named HEM, the authors report that the resulting data structure is 21% larger than the one built with plain MWHC, and lookups are 30–50% slower. A similar overhead was confirmed in our experiments, which are discussed in Section 5.

4 Cache-oblivious peeling

In this section we describe a cache-oblivious algorithm to peel an r-hypergraph. We describe the algorithm for 3-hypergraphs, but it is easy to generalize it to arbitrary r.

4.1 Maintaining incidence lists

To represent the hypergraph throughout the execution of the algorithm, we need a data structure to store the *incidence list* of every vertex v_0, i.e., the list $L_{v_0} = \{(v_0, v_1^0, v_2^0), \ldots, (v_0, v_1^{d-1}, v_2^{d-1})\}$ of *valid* oriented edges whose first vertex is v_0. To realize the peeling algorithm, it is sufficient to implement the following operations on the lists.

- $\mathrm{Degree}(L_{v_0})$ returns the number of edges d in the incidence list of v_0;
- $\mathrm{AddEdge}(L_{v_0}, e)$ adds the edge e to the incidence list of v_0;
- $\mathrm{DeleteEdge}(L_{v_0}, e)$ deletes the edge e from the incidence list of v_0;

- $\mathrm{RetrieveEdge}(L_{v_0})$ returns the only edge in the list if $\mathrm{Degree}(L_{v_0}) = 1$.

For all the operations above, it is assumed that the edge is given through a *valid* orientation. Under this set of operations, the data structure does not need to *store* the actual list of edges: it is sufficient to store a tuple $(v_0, d, \tilde{v}_1, \tilde{v}_2)$, where d is the number of edges, $\tilde{v}_1 = \bigoplus_{j<d} v_1^j$, and $\tilde{v}_2 = \bigoplus_{j<d} v_2^j$, that is, all the vertices of the list in the same position are XORed together. The operations $\mathrm{AddEdge}$ and $\mathrm{DeleteEdge}$ on an edge (v_0, v_1', v_2') simply XOR v_1' into $\tilde{v}_1$ and v_2' into $\tilde{v}_2$, and respectively increment or decrement d. Since all the edges are assumed valid (i.e., it holds that $v_1' < v_2'$) these operations maintain the invariant. When $d = 1$, clearly $\tilde{v}_1 = v_1$ and $\tilde{v}_2 = v_2$ where (v_0, v_1, v_2) is the only edge in L_{v_0}, so it can be returned by $\mathrm{RetrieveEdge}$. If necessary, the data structure can be trivially extended to *labeled* edges (v_0, v_1, v_2, ℓ) by XORing together the labels ℓ into a new field $\tilde{\ell}$. We call this data structure *packed incidence list*, and we refer to this technique as the *XOR trick*. The advantage with respect to maintaining an explicit list, besides the obvious space savings, is that it is sufficient to maintain a single fixed-size record per vertex, regardless of the number of incident edges. This will make the peeling algorithm in the next section substantially simpler and faster. The same trick can be applied to the standard linear-time algorithm, replacing the linked lists traditionally used. As we will see in Section 5, the improvements are significant in both working space and running time.

4.2 Layered peeling

The peeling procedure we present is an adaptation of the CORE procedure presented by Molloy [20]. The basic idea is to proceed in rounds: at each round, all the vertices of degree 1 are removed, and then the next round is performed on the induced subgraph, until either a 2-core is left, or the graph is empty. In the latter case, the algorithm partitions the edges into a sequence of *layers*, one per round, by defining each layer as the set of edges removed in its round. It is easy to see that by concatenating the layers the resulting edge order is a peeling order, regardless of the order within each layer.

The layered peeling process terminates in a small number of rounds: Jiang et al. [14] proved that if the hypergraph is generated randomly with a sparsity above the peeling threshold, then w.h.p. the number of rounds is bounded by $O(\log \log n)$. Moreover, the fraction of vertices remaining in each round decreases double-exponentially. In the following we show how to implement the algorithm in an I/O-efficient way by putting special care in the hypergraph representation and the update step.

Hypergraph representation At each round i, the hypergraph is represented by a list E_i of tuples $(v_0, d, \tilde{v}_1, \tilde{v}_2)$ as described in Section 4.1; each tuple represents the incidence list of v_0. Each list E_i is sorted by its first component v_0. Note that each edge $e = \{v_0, v_1, v_2\}$ needs to be in the incidence list of all its vertices; hence, all the three orientations of e are present in the list E_i.

Construction of E_0 To construct E_0, the edge list for the first round, we put together in a list all the valid orientations of all the edges in the hypergraph. The list is then sorted by v_0, and from the sorted list we can construct the sorted list of incidence lists E_0: after grouping the oriented edges by v_0, we start with the empty packed incidence list $(v_0, 0, 0, 0)$ and, after performing $\mathrm{AddEdge}$ with all the edges in the group, we append it to E_0. The I/O complexity is $O(\mathrm{sort}(n) + \mathrm{scan}(n)) = O(\mathrm{sort}(n))$.

Round update At the beginning of each round we are given the list E_i of edges that are alive at round i, and we produce E_{i+1}. We first scan E_i to find all the tuples L such that $\mathrm{Degree}(L) = 1$; for each tuple, we perform $\mathrm{RetrieveEdge}$ and put the edge in a list D_i, which represents all the edges to be removed in the current round i. The same edge may occur multiple times in D_i under different orientations (if more than one of its vertices have degree 1 in the current round); to remove the duplicates, we sort the oriented edges by their canonical orientation, keep one orientation for each edge, and store them in a list P_i.

Now we need to remove the edges from the hypergraph. To do so, we generate a *degree update list* U_i that contains all the three orientations for each edge in P_i, and sort U_i by v_0. Since both E_i and U_i are sorted by v_0, we can scan them both simultaneously joining them by v_0; for each tuple L_{v_0} in E_i, if no oriented edge starting with v_0 is in U_i the tuple is copied to E_{i+1}, otherwise for each such oriented edge e, $\mathrm{DeleteEdge}(L_{v_0}, e)$ is called to obtain a new L'_{v_0} which is written to E_{i+1} if non-empty. Note that E_{i+1} remains sorted by v_0.

For each round, we scan E_i twice and U_i once, and sort D_i and U_i. The number of I/Os is then $2 \cdot \mathrm{scan}(|E_i|) + \mathrm{scan}(|U_i|) + \mathrm{sort}(|D_i|) + \mathrm{sort}(|U_i|)$. Summing over all rounds, we have $\sum_i (2 \cdot \mathrm{scan}(|E_i|) + \mathrm{sort}(|D_i|) + \mathrm{sort}(|U_i|) + \mathrm{scan}(|U_i|)) = O(\mathrm{sort}(n))$ because each edge belongs to at most three lists D_i and three lists U_i. Since the fraction of vertices remaining at each round decreases doubly exponentially and, thanks to the XOR trick, E_i has exactly one tuple for each vertex alive in the i-th round, the cost of scanning the lists E_i sums up to $O(\mathrm{scan}(n))$ I/Os. Hence, overall the algorithm takes w.h.p. $O(n \log n)$ time and $O(\mathrm{sort}(n))$ I/Os.

4.3 Implementation details

We report here the most important optimizations we used in our implementation.

File I/O Instead of managing file I/Os directly, we use a memory-mapped file by employing a C++ allocator that creates a file-backed area of memory. This way we can use the standard STL containers such as `std::vector` as if they resided in internal memory. We use `madvise` to instruct the kernel to optimize the mapped region for sequential access.

Sorting Our sorting implementation performs two steps: in the first step we divide the domain of the values into k evenly spaced buckets, scanning the array to find the number of values that belong in each bucket, and then moving each value to its own bucket. In the second step, each bucket is sorted using `sort` of the C++ standard library (the procedure is entirely I/O bound, so using, e.g., radix sort has no impact). The number of buckets is chosen so that with w.h.p. each bucket fits in internal memory; since the graph is random, its edges are uniformly distributed, which makes uniform bucketing w.h.p. balanced. To distribute the values into the k buckets, we use a buffer of size T for each bucket; when the buffer is full, it is flushed to disk. Note that this algorithm is technically not cache-oblivious, since it works as long as the available memory M is at least kT; choosing k to be $\Theta(S/M)$, where S is the size of the data to be sorted, requires that M be $\Omega(\sqrt{TS})$. In our implementation we use $T \approx 1$MiB, thus for example $M = 1$GiB is sufficient to sort $\approx$1TiB of data. When this condition holds, the algorithm performs just three scans of the array and it is extremely efficient in practice. Furthermore, contrary to existing cache-oblivious sorting implementations, it is *in-place*, using no extra disk space.

Reusing memory The algorithm as described in Section 4.2 uses a different list E_i for each

round. Since tuples are appended to E_{i+1} at a slower pace than they are read from E_i, we can reuse the same array. A similar trick can be applied to D_i and U_i. Overall, we need to allocate just one array of γn packed incidence lists, and one for the $3n$ oriented edges.

Lists compression Reducing the size of the on-disk data structures can significantly improve I/O efficiency, and hence the running time of the algorithm. The two data structures that take nearly all the space are the lists of packed incidence lists E_i and the lists of edges P_i. Since the lists are read and written sequentially, we can *(de)compress* them on the fly.

Recall that the elements of E_i are tuples of the form $(v_0, d, \tilde{v}_1, \tilde{v}_2)$ sorted by v_0. The first components v_0 of these tuples are gap-encoded with Elias γ codes. The overall size of the encoding is $\sum_{k=1}^{|E_i|}(2\lfloor \log g_k \rfloor + 1)$ bits, where g_k is the k-th gap. Since the gaps sum up to γn, by Jensen's inequality the sum is maximized when the gaps g_k are all equal to $\frac{\gamma n}{|E_i|}$ giving a space bound of $2|E_i|(\log \frac{\gamma n}{|E_i|} + 1)$ bits. Furthermore, this space bound is always at most $2\gamma n$ bits because it is maximized when E_i has size $\gamma n/2$. The degrees d are encoded instead with unary codes; since the sum of the degrees is $3n_i$, where n_i is the number of edges alive in round i, the overall size of their encoding is always upper bounded by $3n$ bits. The other two components, as well as the nodes in P_i, are represented with a fixed-length encoding using $\lceil \log \gamma n \rceil$ bits each. With $\gamma = 1.23$, the overall disk usage is approximately $(5.46 + 11.46\lceil \log \gamma n \rceil)n$ bits. On our largest inputs, using compression instead of plain 64-bit words makes the overall algorithm run about 2.5 times faster.

5 Experimental analysis

Although our code can be easily extended to construct *any* given static function, to evaluate experimentally the performance of the peeling algorithm we tested it on the task of constructing a minimal perfect hash function, as discussed in Section 3. In this task, the peeling process largely dominates the running time.

Testing details The tests of MPHF construction were performed on an Intel Xeon i7 E5520 (Nehalem) at 2.27GHz with 32GiB of RAM, running Linux 3.5.0 x86-64. The storage device is a 3TB Western Digital WD30EFRX hard drive. Before running each test, the kernel page cache was cleared to ensure that all the data were read from disk. The experiments were written in C++11 and compiled with g++ 4.8.1 at `-O3`. We tested the following algorithms.

- Cache-Oblivious: The cache-oblivious algorithm described in Section 4.
- Standard+XOR: The standard linear-time peeling implemented using the packed incidence list, with the purpose of evaluating the impact of the XOR-trick by itself.
- cmph: A publicly available library for minimal perfect hashingin-memory peeling algorithm.

Datasets We tested the above algorithms on the following datasets.

- URLs: a set of ≈ 4.8 billion URLs from the ClueWeb09 dataset[3] (average string length ≈ 67 bytes, summing up to ≈ 304GiB);
- ngrams: a set of ≈ 7.6 billion $\{1, 2, 3\}$-grams obtained from the Google Books Ngrams English dataset[4] (average string length ≈ 23 bytes, summing up to ≈ 168GiB).

[3] `http://lemurproject.org/clueweb09/`

[4] `http://storage.googleapis.com/books/ngrams/books/datasetsv2.html`.

Since the strings are hashed in the first place, the nature of the data is fairly irrelevant: the only aspect that may be relevant is the average string length (that affects the time to load the input from disk). In fact tests on randomly generated data produced the same results.

Experimental results The running time of the algorithms as the number of keys increases is plotted in Figure 1; to evaluate the performance in the regime where the working space fits in main memory, the figure also shows an enlarged version of the first part of the plot.

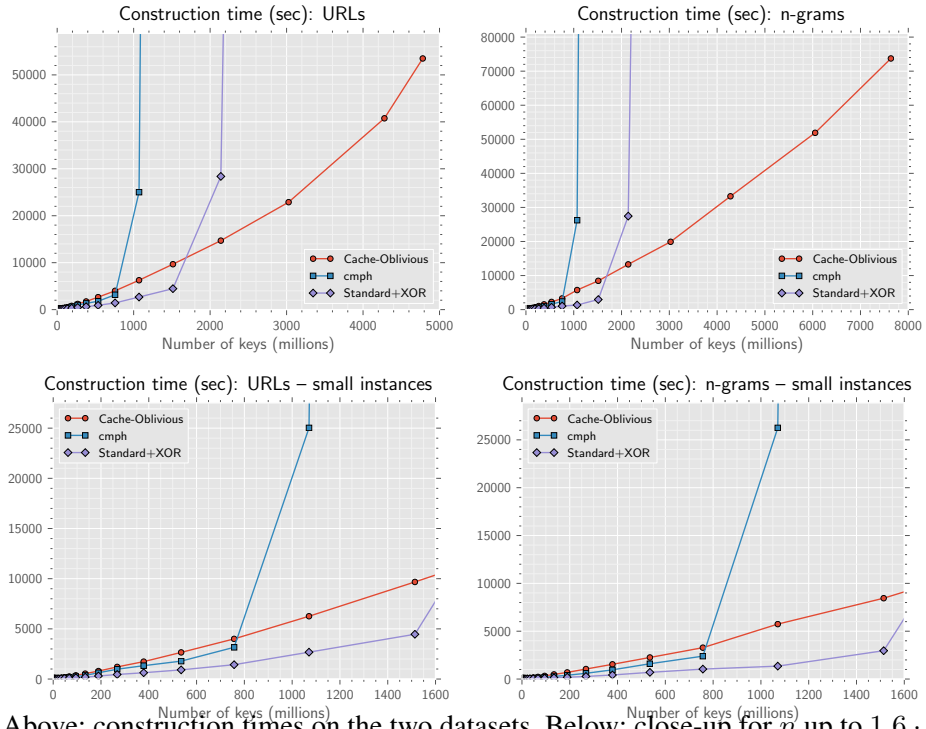

Figure 1: Above: construction times on the two datasets. Below: close-up for n up to $1.6 \cdot 10^9$ keys.

The first interesting observation is that the cache-oblivious algorithm performs almost as well as cpmh, with Cache-Oblivious being slightly slower because it has to perform file I/O even when the working space would fit in memory.

We can also see that the XOR trick pays off, as shown by the performance of Standard+XOR, which is up to 3 times faster than cmph, and the smaller space usage enables to process up to almost twice the number of keys for the given memory budget. Both non-external algorithms, though, cease to be useful as soon as the available memory gets exhausted: the machine, then, starts to thrash because of the random patterns of access to the swap (we killed the processes after 48 hours). Actually, one can make a quite precise estimate of when this is going to happen: cpmh occupies 34.62 bytes/key, as estimated by the authors, whereas Standard+XOR occupies about 26.76 bytes/key, and these figures almost exactly justify the two points where the construction slows down and then stalls. On the other hand, Cache-Oblivious scales well with the input size, exhibiting eventually almost linear performance in our larger input ngrams, while remaining competitive even on small key sets.

Comparison with HEM Finally, we compare our algorithm with HEM [6]. Recall that their technique consists in splitting the set of keys into several buckets and building a separate MPHF for every bucket; at query time, a first-level index is used to drive the query to the correct bucket. Choosing a sufficiently small size for the buckets allows the use of a

	URLs ($4.8 \cdot 10^9$ keys)		ngrams ($7.6 \cdot 10^9$ keys)	
	Lookup	Space	Lookup	Space
	Intel i7			
MWHC	414 ns $\pm$ 0.3%	2.61 b/key	405 ns $\pm$ 0.3%	2.61 b/key
HEM	506 ns $\pm$ 0.3%	3.31 b/key	480 ns $\pm$ 0.3%	3.05 b/key
	AMD Opteron			
MWHC	398 ns $\pm$ 0.1%	2.61 b/key	366 ns $\pm$ 0.1%	2.61 b/key
HEM	475 ns $\pm$ 0.1%	3.31 b/key	441 ns $\pm$ 0.1%	3.05 b/key

Table 1: Space and lookup-time comparison (with relative standard deviation) of HEM and MWHC.

standard internal memory algorithm to construct the bucket MPHF. Although technically not a peeling algorithm, this external-memory solution is simple and elegant.

To make a fair comparison, we re-implemented the HEM algorithm using our sort implementation for the initial bucketing, and the Standard+XOR algorithm to build the bucket MPHFs. The signature function is the same 96-bit hash function used in [6] (which suffice for sets of up to 2^{48} keys), but we employed 64-bit bucket offsets in place of 32-bit, since our key sets are larger than $2^{32}/\gamma$.

The result, as shown in Figure 2, is a construction time between 2 and 6 times smaller than Cache-Oblivious. However, this efficiency has a cost in term of lookup time (because of the double indirection) and size (because of the extra space needed for the first-level index). Since, in most applications, MPHFs are built rarely and queried frequently, the shorter construction time may not be worth the increase in space and query time.

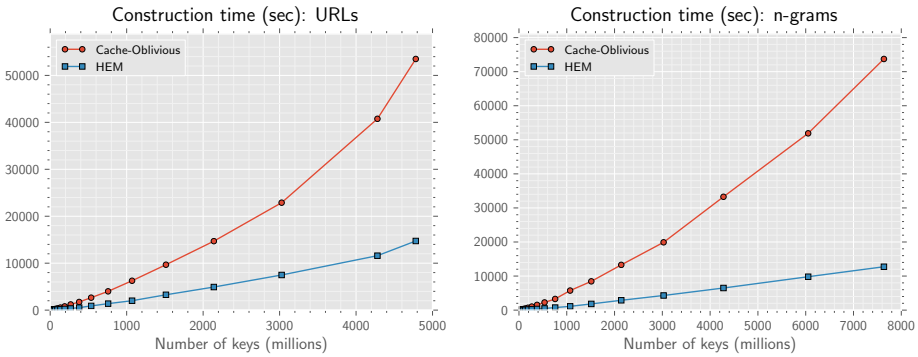

Figure 2: Construction time with the Cache-Oblivious algorithm and HEM [6].

Indeed, as shown in Table 1, the space loss is 17% to 27%. The variability in space overhead is due to the fact that in HEM the number of buckets must be a power of 2, hence the actual average bucket size can vary by a factor of 2 depending on the number of keys.

The evaluation of lookup efficiency is much subtler, as it depends on a number of factors, some of which are subject to hardware architecture. For this reason, we decided to perform the experiments on two quite different machines: the same Intel i7 machine used for the construction experiments (see above) and an AMD Opteron 6276 2.3GHz.

For both machines and both datasets we performed lookups of $10\,$M distinct keys, repeated 10 times. Since lookup times are less than a microsecond, it is impossible to measure individual lookups accurately; for this reason, we divided the lookups into $1{,}525$

batches of 2^{16} keys each, and measured the average lookup time for each batch. Out of these average times, we computed the global average and the standard deviation. The results in Table 1 show that HEM is 18% to 22% slower in all cases. The standard deviation is remarkably small ($\leq 0.3\%$), making the comparison statistically significant.

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