# Zigzag Persistent Homology in Matrix Multiplication Time 

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#### Abstract

We present a new algorithm for computing zigzag persistent homology, an algebraic structure which encodes changes to homology groups of a simplicial complex over a sequence of simplex additions and deletions. Provided that there is an algorithm that multiplies two $n \times n$ matrices in $M(n)$ time, our algorithm runs in $O\left(M(n)+n^{2} \log ^{2} n\right)$ time for a sequence of $n$ additions and deletions. In particular, the running time is $O\left(n^{2.376}\right)$, by result of Coppersmith and Winograd. The fastest previously known algorithm for this problem takes $O\left(n^{3}\right)$ time in the worst case.


## Categories and Subject Descriptors

F.2.2 [Analysis of Algorithms and Problem Complexity]: Nonnumerical Algorithms and Problems; G.2.1 [Discrete Mathematics]: Combinatorics

## General Terms

Algorithms, Theory

## Keywords

Zigzag persistent homology, matrix multiplication.

## 1. INTRODUCTION

Motivation. Since its introduction a decade ago, persistent homology [1] has become an important tool in such wideranging domains as computational biology, geometric processing, machine learning, scientific visualization, and sensor networks. Its success rests on two pillars: a solid theoretical foundation [2], including the celebrated stability result [3], and the availability of fast algorithms [4].

A more recent development is zigzag persistence [5], which is a generalization of ordinary persistence built on algebraic insights. Together with an efficient algorithm in the homological setting, zigzag persistence has already resolved open

[^0]questions in the theoretical study of persistence [6]. In addition to the algebraic generality, it brings an appealing property for algorithm design: one is not constrained to study growing families of spaces as in ordinary persistence, but instead is free to choose whether to grow or shrink the space in question on demand. This flexibility has led to a technique for computing ordinary persistent homology of a real-valued function using space that depends only on the size of the largest levelset, rather than the entire domain [6].

Despite the growing interest in persistence, the complexity of its computation is not well understood. The best currently known algorithms for ordinary and zigzag persistence run in time $\mathrm{O}(M(n))$ and $\mathrm{O}\left(n^{3}\right)$, respectively, in the worst case, where $M(n)$ is the time to multiply two $n \times n$ matrices. In this paper, we present an algorithm that computes zigzag persistent homology (and therefore also ordinary persistent homology) over a finite field in time $\mathrm{O}\left(M(n)+n^{2} \log ^{2} n\right)$. It is not known if any of these algorithms are optimal, and we leave this question open.

We also do not consider the problem of computing homology over infinite fields. In this case it cannot be assumed that arithmetic operations take constant time, but it is known that standard homology can still be computed in $\mathrm{O}(M(n))$ time [7].

Related Work. The computation of homology through Gaussian elimination has been known for some time $[8,9]$. This algorithm has the worst case running time of $\mathrm{O}\left(n^{3}\right)$. Persistent (and standard) homology is essentially Gaussian elimination with known column order and row pivoting. Therefore, it can be computed in time $\mathrm{O}(M(n)$ ) (see [10] for details) over finite fields, using the algorithm for PLU factorization of Bunch and Hopcroft [11], with minor modifications. In the case of zigzag persistent homology, neither row nor column order is known in advance, i.e. both row and column pivoting is required. Some elimination steps are specified by column, others by row, but never both. Furthermore, the two kinds of elimination steps can be arbitrarily interleaved. Due to these complications, it is not clear how to apply [11] and other similar ideas $[12,13]$ to this problem.

The first sub-cubic algorithm to appear incrementally computes the Betti numbers of subcomplexes of triangulations of the three sphere $\mathbb{S}^{3}[14]$. The running time is $n \alpha(n)$, where $n$ is the number of simplices and $\alpha(\cdot)$ is the inverse Ackermann function. An approach to computing Betti numbers using combinatorial Laplacians appears in [15]. It uses the power method on the Laplacian matrix, hence operating via matrix-vector multiplication. However, the number of mul-
tiplications depends on the eigenvalues of the matrix and hence is not easily bounded.

The introduction of persistent homology [1, 2], zigzag persistence $[5,6]$, and other variants $[16,17]$ came with algorithms based on the sequential Gaussian elimination and so had a running time of $O\left(n^{3}\right)$. In [18], an example complex is given where the sequential persistence algorithm takes cubic time. However, in practice, experimental results show close to linear behavior, most often attributed to the sparsity of the involved matrices.

In the literature, most work to speed up homology computation has had a different flavor than our approach, with efforts being made to reduce the size of the input complex using combinatorial operations which preserve the homology [19], most often simplicial collapses. In certain cases, notably 2 -manifolds, it is possible to create an optimal order of collapses to obtain the homology [20]. In the end of this procedure we end up only with critical cells, allowing us to simply read off the homology. However, it was also shown that in the general case, there may not exist a way to collapse the complex to the minimal number of cells [21]. Furthermore, it was shown to be NP-hard to find an order which will result in the minimum number of cells [22]. More recently, a method [23] was given to reduce the size of the complex in the special case of clique complexes. Such complexes are completely determined by their graphs, and the work tries to extract a minimal representation based on maximal cliques. As with the collapsing techniques, it does not come with any guarantees on the size of the output complex.

In this work we use fast matrix multiplication as a blackbox, but we briefly recount the work which began with the celebrated result of Strassen [24], showing that the matrix inversion can be done with an exponent of $\omega \approx 2.8$ rather than 3. This has been followed by numerous improvements, with the currently best known algorithm of CoppersmithWinograd [25]. The current best bound for the exponent is $\omega=2.376$. Finally, we note that this is still an active area of research, where [26] proves that if a group with certain properties exists, so must a quadratic time algorithm for matrix multiplication.

Outline. We review the necessary background in Section 2. We recast the zigzag persistent homology algorithm of [6] in terms of matrix multiplications in Section 3. By batching some of the operations together in Section 4, we obtain the claimed running time.

## 2. BACKGROUND

In this section, we recount the necessary mathematical and algorithmic background. Since the emphasis of this paper is algorithmic, we begin with a simplicial complex $\mathbb{X}$ rather than the usual topological space. Furthermore, everything will be done with simplicial homology over a field $\mathbb{F}$. While mathematically it is often more convenient to work in singular homology, in practice, we always compute in a combinatorial setting.

We now give a brief overview of homology and zigzag persistence. We refer the reader to Munkres [8] or Hatcher [27] for background on homology, Edelsbrunner and Harer [4] for persistence, and Carlsson and de Silva [5] for zigzag persistence. Our primary object is a simplicial complex, which is a set of simplices such that all faces of a simplex are in the
complex and the intersection of two simplices is a (possibly empty) face of both.

We define the chain group $\mathrm{C}_{p}$ as an Abelian group on the set of oriented $p$-simplices in the simplicial complex. A $p$ chain is a linear combination of simplices with coefficients in a group. We restrict ourselves to the case where the coefficients lie in a field, therefore each $\mathrm{C}_{p}$ is a vector space. We also define the boundary operator $\partial_{p}: \mathrm{C}_{p} \rightarrow \mathrm{C}_{p-1}$. In a simplicial complex, the boundary operator is defined on a simplex $\sigma$ as $\partial_{p} \sigma=\sum_{i}(-1)^{i}\left[v_{0}, v_{1}, \ldots, \hat{v}_{i}, \ldots, v_{p}\right]$, where $\hat{v}_{i}$ is deleted from the sequence. Since the operator is linear, the boundary of a chain is the linear combination of the boundaries of the simplices. The boundary operator connects the chain groups into the chain complex: $\ldots \rightarrow$ $\mathrm{C}_{p+1} \xrightarrow{\partial_{p+1}} \mathrm{C}_{p} \xrightarrow{\partial_{p}} \mathrm{C}_{p-1} \rightarrow \ldots$ The boundary operator is a map between vector spaces and is most naturally represented as a matrix where the rows represent $(p-1)$-simplices and the columns represent $p$-simplices.

To define homology, we require two subgroups: the cycle group Z and the boundary group B . The cycle group $\mathrm{Z}_{p}$ is the kernel of the $\partial_{p}$, which is the null space of the boundary matrix. The boundary group $\mathrm{B}_{p}$ is the image of $\partial_{p+1}$. The homology is the quotient group of the two subspaces: $\mathrm{H}_{p}=$ $\mathrm{Z}_{p} / \mathrm{B}_{p}$. By the property that $\partial_{p} \partial_{p+1}=0$, it is not difficult to see that $\mathrm{B}_{p} \subseteq \mathrm{Z}_{p} \subseteq \mathrm{C}_{p}$, meaning the above quotient is well defined. Since these are vector spaces, standard Gaussian elimination can be used to find a basis for each space.

Persistent homology and its recently-introduced generalization, zigzag persistence, are two powerful extensions of the classical homology theory. Instead of working with a fixed simplicial complex, both study a parameterized family of complexes and examine the evolution of homology classes in the induced sequences of homology groups. The study is motivated by data that simultaneously contains features at multiple scales - a common behavior in practice - and, therefore, does not justify a single choice of the scale parameter. In contrast, by examining a data set across all scales, we extract information about meaningful parameter values as well as detect the prominent features in the input.

The starting point of zigzag persistent homology is a sequence of spaces connected with maps

$$
\begin{equation*}
\mathbb{X}_{1} \leftrightarrow \mathbb{X}_{2} \leftrightarrow \ldots \leftrightarrow \mathbb{X}_{n} \tag{1}
\end{equation*}
$$

The above notation means that the map between spaces can point in either direction. For each space we can form a chain complex; the maps between topological spaces induce maps between the chain complexes, $C\left(\mathbb{X}_{1}\right) \leftrightarrow C\left(\mathbb{X}_{2}\right) \leftrightarrow$ $\ldots \leftrightarrow C\left(\mathbb{X}_{n}\right)$. Finally, by passing to homology, we obtain a sequence of vector spaces connected by homomorphisms,

$$
\begin{equation*}
\mathrm{H}\left(\mathbb{X}_{1}\right) \leftrightarrow \mathrm{H}\left(\mathbb{X}_{2}\right) \leftrightarrow \ldots \leftrightarrow \mathrm{H}\left(\mathbb{X}_{n}\right) \tag{2}
\end{equation*}
$$

This sequence is a zigzag module, denoted by $\mathbb{V}$. As shown in [5], $\mathbb{V}$ has a unique decomposition into a direct sum of interval modules, $\mathbb{V}=\bigoplus \mathbb{I}_{[b, d]}$. Each interval module $\mathbb{I}_{[b, d]}$ represents a homology class which exists in all the spaces from $\mathrm{H}\left(\mathbb{X}_{b}\right)$ to $\mathrm{H}\left(\mathbb{X}_{d}\right)$ inclusive. Therefore, we say this class persists from $b$ to $d$. Note that persistent homology is a special case of zigzag persistence where all the maps point one way.

To compute this decomposition, we require compatible bases in all of the individual homology groups and track how they change as we apply the maps. Fortunately, a single
filtration encodes all the required information (see [5] for details).
Definition. The right filtration of a space $\mathbf{H}\left(\mathbb{X}_{i}\right)$ is the collection of its subspaces taking the form

$$
R^{i}=\left(R_{0}, R_{1}, R_{2}, \ldots, R_{i}\right)
$$

satisfying inclusion relations $R_{k} \subseteq R_{l}$ for $k \leq l$. The filtration is then defined inductively. $\overline{R^{0}}=(0)$. If the map points forward: $\mathrm{H}\left(\mathbb{X}_{i}\right) \xrightarrow{f} \mathrm{H}\left(\mathbb{X}_{i+1}\right)$, then the filtration is updated with the images of the map,

$$
R^{i+1}=\left(f\left(R_{0}\right), f\left(R_{1}\right), \ldots, f\left(R_{i}\right), \mathrm{H}\left(\mathbb{X}_{i+1}\right)\right)
$$

If the map points in the other direction: $\mathrm{H}\left(\mathbb{X}_{i}\right) \stackrel{g}{\leftarrow} \mathrm{H}\left(\mathbb{X}_{i+1}\right)$, then the preimages are added,

$$
R^{i+1}=\left(0, g^{-1}\left(R_{0}\right), g^{-1}\left(R_{1}\right), \ldots, g^{-1}\left(R_{i}\right)\right)
$$

By keeping track of the changes in the right filtration as we process the zigzag in (1), we obtain the interval decomposition. In the algorithmic setting, without loss of generality, we can assume that in the sequence of simplicial complexes (1) consecutive complexes differ by a single simplex, i.e. either $\mathbb{X}_{i+1}=\mathbb{X}_{i} \cup \sigma_{i}$, or $\mathbb{X}_{i}=\mathbb{X}_{i+1} \cup \sigma_{i}$.

## 3. SEQUENTIAL ALGORITHM

In this section we express the algorithm of [6] in terms of matrix update operations.

Representation. We start with a sequence of simplicial complexes (1), where consecutive spaces differ by a single simplex. Enumerating all the simplices $\sigma_{1}, \ldots, \sigma_{k}$ that appear in the sequence, we ignore multiplicities: if the same simplex enters, leaves, and then re-enters, we treat it as two simplices in the enumeration. We associate to each simplex $\sigma_{i}$ its interval $\mathbb{I}_{\left[b_{i}, d_{i}\right]}$ in the chain complex zigzag, i.e. $\sigma_{i} \in \mathbb{X}_{j}$ iff $j \in\left[b_{i}, d_{i}\right]$.

As in [6], we maintain the right filtration of the zigzag (2), and represent it after $j-1$ steps via the matrices $Z^{j}, B^{j}$ as well as the matrix $C^{j}$, which maintains the bounding chains. Denoting the boundary matrix of the simplicial complex with $D$, we have $0=D Z^{j}, Z^{j} B^{j}=D C^{j}$. Furthermore, the right filtration of the group $\mathrm{H}\left(\mathbb{X}_{j}\right)=\left(R_{0}^{j}, R_{1}^{j}, \ldots\right)$ is represented via

$$
R_{k}^{j}=\operatorname{span}\left(\left\{z+\mathrm{B} \mid z \in \mathrm{Z}_{k}^{j} \text { and } \mathrm{B}=\operatorname{span}\left(Z^{j} B^{j}\right)\right\}\right),
$$

where $\mathbf{Z}_{k}^{j}$ denotes the subset of the cycle group spanned by the first $k$ columns of the matrix $Z^{j}$. We also implicitly maintain the birth vector associated to the right filtration, which allows us to output the persistence intervals; however, this detail is irrelevant to our argument, and we ignore it from now on.

In addition to the matrices $D, Z^{j}, C^{j}, B^{j}$ of [6], we introduce matrices $E^{j}, F^{j}$. The matrices $E^{j}$ and $F^{j}$ keep track of the "past" and "future" simplices, respectively. Their columns correspond to the simplices that have been removed in the case of $E^{j}$, or not yet added in the case of $F^{j}$. Their boundaries are expressed using the auxiliary matrices $G^{j}, H^{j}, K^{j}, L^{j}, R^{j}, S^{j}$, as defined by Equation (3) - for instance, $Z_{p-1}^{j} H_{p}^{j}$ is the contribution of the existing cycles to the boundaries of the "future" simplices, fully expressed as,

$$
D_{p} F_{p}=Z_{p-1} H_{p}+C_{p-1} K_{p}+F_{p-1} L_{p} .
$$

Similarly, for the boundaries of the "past" simplices,

$$
D_{p} E_{p}=E_{p-1} G_{p}+Z_{p-1} R_{p}+C_{p-1} S_{p}
$$

We say that the matrix $H^{j}$ is independent of the matrix $B^{j}$ if any non-zero row in matrix $B^{j}$ is zero in matrix $H^{j}$.

Sequential Invariant. After step $j-1$ of the zigzag, the following conditions must hold:
1.
$D_{p}\left[\begin{array}{c|c|c|c}E_{p}^{j} & 0 & 0 & 0 \\ \hline 0 & Z_{p}^{j} & C_{p}^{j} & F_{p}^{j}\end{array}\right]=$
$\left[\begin{array}{c|c|c|c}E_{p-1}^{j} & 0 & 0 & 0 \\ \hline 0 & Z_{p-1}^{j} & C_{p-1}^{j} & F_{p-1}^{j}\end{array}\right]\left[\begin{array}{c|c|c|c}G_{p}^{j} & 0 & 0 & 0 \\ \hline R_{p}^{j} & 0 & B_{p}^{j} & H_{p}^{j} \\ \hline S_{p}^{j} & 0 & 0 & K_{p}^{j} \\ \hline 0 & 0 & 0 & L_{p}^{j}\end{array}\right]$
For convenience, we abbreviate the matrices in (3) as $D \Phi_{p}^{j}=\Phi_{p-1}^{j} \Gamma_{p}^{j}$.
2. Matrix $B_{p}^{j}$ has exactly one non-zero element per column, and at most one per row. Matrix $H_{p}^{j}$ is independent of $B_{p}^{j}$.
3. The columns of matrix $F_{p}^{j}$ are linearly independent. Matrix $Z_{p}^{j}$ forms a basis for the $p$-cycles $\mathbf{Z}_{p}\left(\mathbb{X}_{j}\right)$. The boundary $D C_{p}^{j}$ forms a basis for the $(p-1)$-boundaries $\mathrm{B}_{p}\left(\mathbb{X}_{j}\right)$.
4. $R_{k}^{j}=\operatorname{span}\left(\left\{z+\mathrm{B} \mid z \in \mathrm{Z}_{k}^{j}\right.\right.$ and $\left.\left.\mathrm{B}=\operatorname{span}\left(Z^{j} B^{j}\right)\right\}\right)$.

Additionally, we choose a particular ordering for the rows and columns of our matrices.

Ordering. The rows of the matrices $Z^{j}, C^{j}, E^{j}$, and $F^{j}$ correspond to the individual simplices in the complex, and we order them by the removal time of those simplices. Consequently, the rows and the columns of the boundary matrix $D$ are also ordered by the removal time. The columns of the matrix $F^{j}$ represent future simplices, and we order them by the addition time. The columns of the matrix $E^{j}$ represent past (removed) cycles and chains, and we order them by the removal time.

From the invariant and ordering requirements it follows that, initially, $L_{p}^{0}$ is the boundary matrix with rows and columns ordered by addition, while $F_{p}^{0}$ is a permutation matrix, its columns and rows are indexed by simplices ordered by addition and removal, respectively. The above ordering is insignificant in the rest of this section, however, it becomes critical in Section 4 when we describe how to divide and conquer the necessary computation.

Four cases. In the remainder of the section, we express the update performed to the matrices after each step as matrix multiplication.
$\left[\begin{array}{c|c|c|c}E_{p}^{j+1} & 0 & 0 & 0 \\ \hline 0 & Z_{p}^{j+1} & C_{p}^{j+1} & F_{p}^{j+1}\end{array}\right]=\left[\begin{array}{c|c|c|c}E_{p}^{j} & 0 & 0 & 0 \\ \hline 0 & Z_{p}^{j} & C_{p}^{j} & F_{p}^{j}\end{array}\right] M_{p}^{j}$

$$
\begin{aligned}
& {\left[\begin{array}{c|c|c|c}
G_{p}^{j+1} & 0 & 0 & 0 \\
\hline R_{p}^{j+1} & 0 & B_{p}^{j+1} & H_{p}^{j+1} \\
\hline S_{p}^{j+1} & 0 & 0 & K_{p}^{j+1} \\
\hline 0 & 0 & 0 & L_{p}^{j+1}
\end{array}\right]=} \\
& \\
& \\
& \\
&
\end{aligned}
$$

Step $j$ of the zigzag can be one of the two possibilities: an addition of a simplex, $\mathbb{X}_{j+1}=\mathbb{X}_{j} \cup \sigma_{i}$, or a removal of a simplex, $\mathbb{X}_{j}=\mathbb{X}_{j+1} \cup \sigma_{i}$. Each of the two cases can have one of two outcomes: a birth or a death of a homology class. Before we describe the four possibilities, we establish an equivalence between the matrices.

Lemma 1 (Forward Step Equivalence Lemma). If the $j$-th step of the zigzag is the addition of the simplex $\sigma_{i}$, then $L_{p}^{j}[i]=0, K_{p}^{j}[i]=0$, and $D F_{p}^{j}[i]=Z_{p-1}^{j} H_{p}^{j}[i]$; in particular, $D F_{p}^{j}[i]=0 \Leftrightarrow H_{p}^{j}[i]=0$.

Proof. By the first condition of the sequential invariant, $D F_{p}=Z_{p-1} H_{p}+C_{p-1} K_{p}+F_{p-1} L_{p}$. The last term, $F_{p-1}^{j} L_{p}^{j}[i]$, is zero when we are adding simplex $\sigma_{i}$, since it represents the boundary of $\sigma_{i}$ among the "future" simplices. The assumption that every step of our zigzag is a homology group of a simplicial complex implies that the boundary of a simplex cannot occur in the future. From the invariant, the columns of the matrix $F_{p-1}^{j}$ are linearly independent, therefore, $L_{p}^{j}[i]=0$. Taking the boundary of the remaining equation (and recalling that the boundary matrix squared is 0 ), we get $0=D D F_{p}^{j}[i]=D Z_{p-1}^{j} H_{p}^{j}[i]+D C_{p-1}^{j} K_{p}^{j}[i]=$ $D C_{p-1}^{j} K_{p}^{j}[i]$, where the last equality follows since the boundary of every cycle in the matrix $Z_{p-1}^{j}$ is zero, by definition. Therefore, $K_{p}^{j}[i]=0$ since $D C_{p-1}^{j}$ is a basis by the third condition of the invariant.

We get $D F_{p}^{j}[i]=Z_{p-1}^{j} H_{p}^{j}[i]$. Since the columns of matrix $Z_{p}^{j}$ are linearly independent, $D F_{p-1}^{j}[i]=0$ iff $H_{p}^{j}[i]=0$.

Birth after addition. This case is characterized by $D F^{j}[i]=0$, or, by Lemma 1 equivalently, $H^{j}[i]=0$. In particular, $F^{j}[i]$ is the newly born cycle (indeed, it contains the new simplex $\sigma_{i}$ ). We append it to the cycle matrix $Z^{j}$. The update matrix for this operation is

$$
M_{p}^{j}=\left[\begin{array}{c|cc|c|c}
I & 0 & 0 & 0 & 0  \tag{4}\\
\hline 0 & I & 0 & 0 & 0 \\
\hline 0 & 0 & 0 & I & 0 \\
\hline 0 & 0 & 1 & 0 & 0 \\
0 & 0 & 0 & 0 & I
\end{array}\right] .
$$

Death after addition. This case is characterized by $D F_{p}^{j}[i] \neq 0$, or, equivalently by Lemma $1, H_{p}^{j}[i] \neq 0$. Moreover, $D F_{p}^{j}[i]=Z_{p-1}^{j} H_{p}^{j}[i]$ represents the cycle that gets killed by the addition of $\sigma_{i}$. Indeed, since $H_{p}^{j}[i]$ is independent of $B_{p}^{j}$ and non-zero, it means that $Z_{p-1}^{j} H_{p}^{j}[i]$ is not a boundary before step $j$. On the other hand, by definition, $D F_{p}^{j}[i]$ is a boundary after the addition of simplex $\sigma_{i}$. Let $a$ be the last non-zero entry in the column $H_{p}^{j}[i]$. We split the column using this entry into $H_{p}^{j}[i]=\left[\begin{array}{lll}\mathbf{h}^{T} & a & \mathbf{0}^{T}\end{array}\right]^{T}$, where
$\mathbf{h}$ is a vector. Suppose $a$ is in the row $k$. We split the row as follows $H_{p}^{j}[k, \cdot]=\left[a \mathbf{g}^{T}\right]$, where $\mathbf{g}$ is a vector. $F_{p}^{j}[i]$ becomes a bounding chain, and therefore moves to $C_{p}^{j}$. Its boundary replaces the column $k$ of $Z_{p-1}^{j}$ (which is a change of basis in $Z_{p-1}^{j}$ ). We insert a column with a single 1 in row $k$ into the matrix $B_{p}^{j}$ to reflect the death of the cycle $Z_{p-1}^{j}[k]$. We use this column to zero out row $k$ of $H_{p}^{j}$, thus keeping $H_{p}^{j}$ independent of $B_{p}^{j}$. The necessary update operations are

$$
\begin{align*}
M_{p-1}^{j} & =\left[\begin{array}{r|rrr|r|c}
I & 0 & 0 & 0 & 0 & 0 \\
\hline 0 & I & \mathbf{h} / a & 0 & 0 & 0 \\
0 & 0 & 1 / a & 0 & 0 & 0 \\
0 & 0 & 0 & I & 0 & 0 \\
\hline 0 & 0 & 0 & 0 & I & 0 \\
\hline 0 & 0 & 0 & 0 & 0 & I
\end{array}\right],  \tag{5}\\
M_{p}^{j} & =\left[\begin{array}{c|c|cc|c}
I & 0 & 0 & 0 & 0 \\
\hline 0 & I & 0 & 0 & 0 \\
\hline 0 & 0 & I & 0 & 0 \\
\hline 0 & 0 & 0 & 1 & -\mathbf{g}^{T} \\
0 & 0 & 0 & 0 & I
\end{array}\right] .
\end{align*}
$$

Remark 1. A subtle, but important consequence of the matrix $H^{j}$ being independent of the matrix $B^{j}$ is that the column $F^{j}[i]$ arrives "ready": we do not need to reduce it at the beginning of an individual step. This fact is crucial for the speed-up explained in the next section.
Removing a simplex $\sigma_{i}$ corresponds to nullifying the rows $i$ in the matrices $Z^{j}, C^{j}$, and $F^{j}$.

Birth after removal. This case is characterized by $Z_{p}^{j}[i, \cdot]=0$, i.e., no cycle contains simplex $\sigma_{i}$. Let $a$ be the rightmost non-zero entry in row $i$ of matrix $C_{p}^{j}$, and suppose it lies in column $k$, we have $C_{p}^{j}[i, \cdot]=\left[\begin{array}{lll}\mathbf{c}^{T} & a & \mathbf{0}^{T}\end{array}\right]$; specifically, $C_{p}^{j}[i, 1 . . k-1]=\mathbf{c}^{T}, C_{p}^{j}[i, k]=a$. Let $B_{\mathbf{c}}^{j}$ be the part of the matrix $B_{p}^{j}$ corresponding to the columns $\mathbf{c}^{T}$ in Equation (3), i.e. $B_{\mathbf{c}}^{j}=B_{p}^{j}[\cdot, 1 . . k]$. Let $\mathbf{f}^{T}$ be the $i$-th row of the matrix $F_{p}^{j}$.
1: The boundary $D C_{p}^{j}[k]=Z_{p-1}^{j} B_{p}^{j}[k]$ is the newly born cycle. We prepend it to the matrix $Z^{j}$. In the operations below, suppose the 1 in the column $B_{p}^{j}[k]$ is in row $l$.
2: We subtract column $C_{p}^{j}[k] / a$ from the other columns in $C_{p}^{j}$ and all columns of $F_{p}^{j}$.
3: To keep track of the "past", we move column $k$ from $C_{p}^{j}$ to $E_{p}^{j}$ (with the corresponding update in $\Gamma_{p}$ ).
4: The term $-\left(\mathbf{c}^{T} / a\right)\left(B_{\mathbf{c}}^{j}\right)^{T}$ below undoes the effect of Step 2 on matrix $B_{p}^{j}$.
The operations are

$$
\begin{align*}
M_{p-1}^{j} & =\left[\begin{array}{c|ccc|c|c}
I & 0 & 0 & 0 & 0 & 0 \\
\hline 0 & 0 & I & 0 & 0 & 0 \\
0 & 1 & -\left(\mathbf{c}^{T} / a\right)\left(B_{\mathbf{c}}^{j}\right)^{T} & 0 & 0 & 0 \\
0 & 0 & 0 & I & 0 & 0 \\
\hline 0 & 0 & 0 & 0 & I & 0 \\
\hline 0 & 0 & 0 & 0 & 0 & I
\end{array}\right] \text { (row } l \text { ) } \\
M_{p}^{j} & =\left[\begin{array}{cc|c|cc|c}
I & 0 & 0 & 0 & 0 & 0 \\
\hline 0 & 0 & I & 0 & 0 & 0 \\
\hline 0 & 0 & 0 & I & 0 & 0 \\
0 & 1 & 0 & -\mathbf{c}^{T} / a & 0 & -\mathbf{f}^{T} / a \\
0 & 0 & 0 & 0 & I & 0 \\
\hline 0 & 0 & 0 & 0 & 0 & I
\end{array}\right] \tag{6}
\end{align*}
$$

Death after removal. This case is characterized by $Z_{p}^{j}[i, \cdot] \neq 0$, i.e. there is a cycle that contains $\sigma_{i}$. Let $a$ be the leftmost non-zero entry in this row, we have $Z_{p}^{j}[i, \cdot]=$ $\left[\begin{array}{lll}\mathbf{0}^{T} & a & \mathbf{z}^{T}\end{array}\right]$. Furthermore, let $C_{p}^{j}[i, \cdot]=\mathbf{c}^{T}$ be the $i$-th row of matrix $C_{p}^{j}$ and let $F_{p}^{j}[i, \cdot]=\mathbf{f}^{T}$ be the $i$-th row of matrix $F_{p}^{j}$. We zero out these rows by adding multiples of the column containing element $a$ to the other columns, while moving that column into the "past" matrix $E^{j}$.

Operations

$$
M_{p}^{j}=\left[\begin{array}{cc|cc|c|c}
I & 0 & 0 & 0 & 0 & 0  \tag{7}\\
\hline 0 & 0 & I & 0 & 0 & 0 \\
0 & 1 & 0 & -\mathbf{z}^{T} / a & -\mathbf{c}^{T} / a & -\mathbf{f}^{T} / a \\
0 & 0 & 0 & I & 0 & 0 \\
\hline 0 & 0 & 0 & 0 & I & 0 \\
\hline 0 & 0 & 0 & 0 & 0 & I
\end{array}\right] .
$$

In both removal cases, the purpose of zeroing out $\mathbf{f}^{T}$ is to satisfy statement 1 of the Invariant, specifically, to make sure that the row of the freshly removed simplex $\sigma_{i}$ is zero in the future matrix $F_{p}^{j}$.

Remark 2. In the addition cases, we only inspect the column of the matrix $H_{p}^{j}$ corresponding to the $j$-th step of the zigzag, both to make a decision about birth or death, and to construct the update matrices. Similarly, in the removal cases we only inspect the rows of the matrices $Z_{p}^{j}$ or $C_{p}^{j}$ corresponding to the $j$-th step of the zigzag.

In the next section, we divide our matrices into smaller blocks, always ensuring that the necessary parts are available to the procedure responsible for the $j$-th step.

Correctness. The above operations are almost the same as in [6]; in Appendix A, we focus on verifying the new parts of the invariant.

## 4. HIERARCHICAL ALGORITHM

In this section we take advantage of the Remarks 1 and 2 above, and construct a divide-and-conquer algorithm for processing a zigzag of simplicial complexes. We emphasize that our algorithm is simply an efficient implementation of the algorithm from Section 3 by changing the order in which various matrix updates are applied.

For simplicity, we assume that the number of steps in the zigzag is a power of two. Throughout this section, $M(n)$ denotes time to multiply two $n \times n$ matrices.

Restrictions. The main idea of the hierarchical algorithm is that the elementary updates $M_{p}^{j}$ are not applied immediately, but piecewise. Each $M^{j}$ is split into submatrices, which are applied at different times to corresponding submatrices of $\Phi^{j}$ and $\Gamma^{j}$. Recall that rows and columns of $M^{j}, \Phi^{j}, \Gamma^{j}$ are associated with the sequence of insertions and removals (cf. Section 3, Ordering). Not surprisingly, the submatrices of interest to the hierarchical algorithm are associated with contiguous subsequences of operations. We refer to these submatrices as restrictions, and we define them as follows.

Let $i \in[a, b+1]$ be a step. Restriction of matrix $\Phi_{p}^{i}$ to steps $[a, b]$, denoted by $\Phi_{p}^{i ;[a, b]}$, is the submatrix of $\Phi_{p}^{i}$ obtained by removing:

- rows corresponding to $p$-simplices removed outside steps $[a, b]$,
- columns corresponding to $p$-simplices added after step $b$, and
- columns corresponding to $p$-cycles/chains removed before step $a$.

In other words (as it follows from Ordering), to get $\Phi_{p}^{i ;[a, b]}$ from $\Phi_{p}^{i}$, we remove a few topmost and bottommost rows, and a few leftmost and rightmost columns.

We define restrictions of other matrices similarly. Restriction of matrix $\Gamma_{p}^{i}$ to steps $[a, b]$, denoted by $\Gamma_{p}^{i ;[a, b]}$, is the submatrix of $\Gamma_{p}^{i}$ obtained by removing rows corresponding to ( $p-1$ )-simplices added after step $b$, rows corresponding to ( $p-1$ )-cycles/chains removed before step $a$, columns corresponding to $p$-simplices added after step $b$, and columns corresponding to $p$-cycles/chains removed before step $a$.

Let $i \in[a, b]$ be a step. Restriction of update matrix $M_{p}^{i}$ to steps $[a, b]$, denoted by $M_{p}^{i ;[a, b]}$, is the submatrix obtained by removing rows and columns corresponding to $p$-simplices added after step $b$, rows and columns corresponding to $p$ cycles/chains removed before step $a$.

Operation tree. We can view the hierarchical algorithm below as an in-order traversal of a binary tree. The leaves of the tree correspond to individual steps of the zigzag (simplex additions and removals). Each tree vertex represents an invocation of H-ZZPH.


For the rest of this section we assume, unless noted otherwise, that v is a generic tree node whose leaf descendants correspond to steps $[a, b]$. Also, we will often use a tree node as a shorthand for its set of leaf descendants. For example, if $i \in[a, b]$, then we can write $i \in \mathrm{v}, \Phi_{p}^{i ; \mathrm{v}}, \Gamma_{p}^{i ; \mathrm{v}}, M_{p}^{i ; \mathrm{v}}$, etc.

The main recursive procedure H-ZZPH appears below. Lemma 2 formally specifies what $\mathrm{H}-\mathrm{ZZPH}(\mathrm{v})$ does. In a nutshell, it solves the problem for a subsequence of steps given by v, with all matrices restricted to v. More precisely, given initial matrices $\Phi_{p}^{a}$ and $\Gamma_{p}^{a}$, it computes final matrices $\Phi_{p}^{b+1}$ and $\Gamma_{p}^{b+1}$, and all the intermediate elementary updates $M_{p}^{i}, i \in \mathrm{v}$, where all the input and output matrices are restricted to v. Since the root of the tree corresponds to all zigzag steps, top level of recursion returns unrestricted matrices, as desired. $\mathrm{H}-\mathrm{ZZPH}$ is implemented as follows.

[^1]Unless v is a leaf, $\mathrm{H}-\mathrm{ZZPH}(\mathrm{v})$ calls itself recursively on the two children of $v$. Generically, the left child of $v$ is l, its right child is $r$, and the number of leaf descendents of 1 and $r$ (i.e., half the number of leaf descendents of v ) is $k$. We denote by $c$ the last step of l, i.e., $c=a+k-1$. Procedure Left-to-Right loosens the restriction of the result of the first recursive call; it computes $M_{p}^{i ; \mathrm{v}}$ from $\left\{M_{p}^{i ; 1}: i \in \mathrm{l}\right\}$. Procedure Right-To-Left does the symmetric operation for the second recursive call; it computes $M_{p}^{i, \mathrm{v}}$ from $\left\{M_{p}^{i, \mathrm{r}}\right.$ : $i \in \mathrm{r}\}$. Procedure Apply-Left simply applies the result of Left-To-Right to $\Phi_{p}^{a ; \mathrm{v}}$ and $\Gamma_{p}^{a ; \mathrm{v}}$ to compute $\Phi_{p}^{c+1 ; \mathrm{v}}$ and $\Gamma_{p}^{c+1 ; \mathrm{v}}$; this establishes the the precondition for the second recursive call.

Structure of matrices. Our algorithm exploits the structure of $\Phi, \Gamma$, and $M$ matrices in order to multiply them quickly. To describe this structure, we introduce a notion of width.

Definition. An $n \times n$ matrix has width $k$, for some $k \leq n$, if it can be written as $P+U V^{T}$, where $U, V$ are $n \times k$ matrices, and $P$ is an $n \times n$ matrix with at most one nonzero entry in any row or column. The triple $(P, U, V)$ is called the width- $k$ representation of the matrix.

In other words, a width- $k$ matrix is a sum of a sparse matrix and a rank- $k$ matrix. The key property is that matrices can be multiplied in time proportional to their width (cf. Lemma 5). We assume that matrices are also stored in their appropriate low-width representation, if they have one.

It is obvious that elementary update matrices $M_{p}^{i}$ have width-1 representation $P+u v^{T}$, where $P$ is a permutation matrix, so that $P^{-1}=P^{T}$ (see Section 3). Moreover, due to the structure of $\Gamma_{p}^{i}$, any restriction of $\Gamma_{p}^{i}$ to $k$ steps has width $O(k)$. Finally, any restriction of $\Phi_{p}^{i}$ to $k$ steps is a $k \times n$ ("skinny") matrix. We use all these facts for efficient computation (cf. Lemma 3, 4, 5).

Notice that width- $k$ representation is not unique. Product of $k$ width- 1 matrices is a width- $k$ matrix (cf. Lemma 5). In this case, two particular width- $k$ representations are of special interest; we define them by showing how to compute them. Let $\left\{\left(P_{i}, u_{i}, v_{i}\right): i=1, \ldots, k\right\}$ be width-1 representations for $k$ width- 1 matrices. The Prefix and Suffix representation of $\prod_{i=1}^{k}\left(P_{i}+u_{i} v_{i}^{T}\right)$ are defined by

```
\(\operatorname{Prefix}\left(P_{i}, u_{i}, v_{i} \mid i=1, \ldots, k\right)\)
    if \(k=1\) then
        return \(P_{1}, u_{1}, v_{1}\)
    \(Q_{1}, X_{1}, Y_{1} \leftarrow \operatorname{Prefix}\left(P_{i}, u_{i}, v_{i} \mid i \leq\lfloor k / 2\rfloor\right)\)
    \(Q_{2}, X_{2}, Y_{2} \leftarrow \operatorname{Prefix}\left(P_{i}, u_{i}, v_{i} \mid i>\lfloor k / 2\rfloor\right)\)
    return \(Q_{1} Q_{2},\left[\begin{array}{ll}X_{1} & \left(Q_{1}+X_{1} Y_{1}^{T}\right) X_{2}\end{array}\right],\left[\begin{array}{ll}Q_{2}^{T} Y_{1} & Y_{2}\end{array}\right]\)
    \(\operatorname{SUFFIX}\left(P_{i}, u_{i}, v_{i} \mid i=1, \ldots, k\right)\)
    if \(k=1\) then
        return \(P_{1}, u_{1}, v_{1}\)
    \(Q_{1}, X_{1}, Y_{1} \leftarrow \operatorname{SuFFix}\left(P_{i}, u_{i}, v_{i} \mid i \leq\lfloor k / 2\rfloor\right)\)
    \(Q_{2}, X_{2}, Y_{2} \leftarrow \operatorname{SuFFIX}\left(P_{i}, u_{i}, v_{i} \mid i>\lfloor k / 2\rfloor\right)\)
    return \(\left.Q_{1} Q_{2},\left[\begin{array}{ll}X_{1} & Q_{1} X_{2}\end{array}\right],\left[\begin{array}{lll}\left(Q_{2}+X_{2} Y_{2}^{T}\right.\end{array}\right)^{T} Y_{1} \quad Y_{2}\right]\)
```

These two forms are significant because from the output of Prefix (resp. Suffix) we can easily compute a low-width representation of any prefix $\prod_{i=1}^{l}\left(P_{i}+u_{i} v_{i}^{T}\right)$ (resp. suffix $\left.\prod_{i=l}^{k}\left(P_{i}+u_{i} v_{i}^{T}\right)\right), 1 \leq l \leq k$, of the product by zeroing out
columns and permuting. This property will be useful in implementing algorithms Left-to-Right and Apply-Left. In particular, we exploit it in equations (8) and (9) below.

To see that the output of the two algorithms has the claimed form, consider the following operation. First, zero out the second half of the second matrix in the output of Prefix. Then, undo the permutation corresponding to the second half of the product by right-multiplying the output expression of Prefix by $Q_{2}^{T}$,

$$
\left(Q_{1} Q_{2}+\left[\begin{array}{ll}
X_{1} & 0
\end{array}\right]\left[Q_{2}^{T} Y_{1} Y_{2}\right]^{T}\right) Q_{2}^{T}=Q_{1}+X_{1} Y_{1}^{T}
$$

The result is precisely the first half of the product. Symmetrically, in SUFFIX, zeroing the first half of the second matrix and left-multiplying with $Q_{1}^{T}$ yields $Q_{2}+X_{2} Y_{2}^{T}$.

Left-to-Right. The goal of this procedure is to extend the restriction of $\left\{M_{p}^{i}: i \in \mathrm{l}\right\}$, from 1 to v . Definition of restrictions of $M_{p}^{i}$ implies that the only rows/columns in $M_{p}^{i ; \mathrm{v}}$ which are not present in $M_{p}^{i ; 1}$ are those corresponding to $p$-simplices added in the steps of r . In other words, $M_{p}^{i ; \mathrm{v}}$ is missing fewer bottommost rows and rightmost columns compared to $M_{p}^{i ; 1}$.

For $i \in 1$, let $M_{p}^{i ; 1}=P_{i}+u_{i} v_{i}^{T}$, hence

$$
M_{p}^{i ; \mathrm{v}}=\left[\begin{array}{cc}
P_{i} & 0 \\
0 & I
\end{array}\right]+\left[\begin{array}{c}
u_{i} \\
0
\end{array}\right]\left[\begin{array}{c}
v_{i} \\
w_{i}
\end{array}\right]^{T}
$$

where the unknown $w_{i}$ expresses how to apply update $i$ to the leaves of r . The Prefix form of $\prod_{i=a}^{c} M_{p}^{i ; 1}$ is

$$
\prod_{i=a}^{c} M_{p}^{i, 1}=\prod_{i=a}^{c} P_{i}+\left[\begin{array}{lll}
u_{a}^{\prime} & \cdots & u_{c}^{\prime}
\end{array}\right]\left[\begin{array}{lll}
v_{a}^{\prime} & \cdots & v_{c}^{\prime}
\end{array}\right]^{T}
$$

The Prefix form of $\prod_{i=a}^{c} M_{p}^{i ; \mathrm{v}}$ is
$\prod_{i=a}^{c} M_{p}^{i ; \mathrm{v}}=\left[\begin{array}{ccc}\prod_{i=a}^{c} P_{i} & 0 \\ 0 & I\end{array}\right]+\left[\begin{array}{ccc}u_{a}^{\prime} & \cdots & u_{c}^{\prime} \\ 0 & \cdots & 0\end{array}\right]\left[\begin{array}{ccc}v_{a}^{\prime} & \cdots & v_{c}^{\prime} \\ w_{a} & \cdots & w_{c}\end{array}\right]^{T}$.
Note that using Prefix is crucial for $w_{a}, \ldots, w_{c}$ to stay unchanged. We will need prefixes $\prod_{j=a}^{i} M_{p}^{i, \mathrm{v}}$ of this product, and we can access them by zeroing out and permuting as claimed above.

$$
\left.\begin{array}{l}
\prod_{j=a}^{i} M_{p}^{j ; \mathrm{v}}=\left[\begin{array}{cccc}
\prod_{j=a}^{i} P_{j} & 0 \\
0 & I
\end{array}\right]+ \\
{\left[\begin{array}{ccccc}
u_{a}^{\prime} & \cdots & u_{i}^{\prime} & 0 & \cdots
\end{array} 0\right.}  \tag{8}\\
0
\end{array} \cdots \cdots \begin{array}{cccc}
0
\end{array}\right]\left[\begin{array}{ccc}
v_{a}^{\prime} & \cdots & v_{c}^{\prime} \\
w_{a} & \cdots & w_{c}
\end{array}\right]^{T}\left[\begin{array}{ccc}
\left(\prod_{j=i+1}^{c} P_{j}\right)^{T} & 0 \\
0 & I
\end{array}\right] .
$$

The algorithm proceeds in the order of increasing dimension $p$. Therefore, we can assume that $M_{p-1}^{i ; \mathrm{v}}=Q_{i}+x_{i} y_{i}^{T}$ has already been computed for all $i \in l$. In the base case $p=0$, we have $M_{p-1}^{i ; \mathrm{v}}=I$, which is already in the required form. The Suffix form of $\left(\prod_{i=a}^{c} M_{p-1}^{i ; \mathrm{v}}\right)^{-1}$ is

$$
\left(\prod_{i=a}^{c} M_{p-1}^{i ; \mathrm{v}}\right)^{-1}=\left(\prod_{i=a}^{c} Q_{i}\right)^{T}+\left[\begin{array}{lll}
x_{a}^{\prime} & \cdots & x_{c}^{\prime}
\end{array}\right]\left[\begin{array}{lll}
y_{a}^{\prime} & \cdots & y_{c}^{\prime}
\end{array}\right]^{T}
$$

We will need suffixes $\left(\prod_{j=a}^{i} M^{j ; \mathrm{v}}\right)^{-1}$ of this product, and we
use the same zero-and-permute strategy to access them.

$$
\begin{align*}
& \left(\prod_{j=a}^{i} M^{j ; \mathrm{v}}\right)^{-1}=\left(\prod_{j=a}^{i} Q_{j}\right)^{T}+  \tag{9}\\
& \quad\left(\prod_{j=i+1}^{c} Q_{j}\right)\left[\begin{array}{ll}
x_{a}^{\prime} \cdots x_{i}^{\prime} & 0 \cdots 0
\end{array}\right]\left[y_{a}^{\prime} \cdots y_{c}^{\prime}\right]^{T}
\end{align*}
$$

To find the unknown $w_{i}$, we express each one as a linear function of $w_{j}, j<i$ : we compute column vectors $\alpha_{i}, \beta_{i}$ such that $w_{i}=\left[\begin{array}{lll}w_{a} & \cdots & w_{c}\end{array}\right] \alpha_{i}+\beta_{i}$, and only the leading $i-a$ entries of $\alpha_{i}$ can be non-zero. We then find unknown vectors $w_{a}, \ldots, w_{c}$ by solving the linear system. It remains to explain how to compute $\alpha_{i}$ and $\beta_{i}$.

First of all, note that $w_{i}=0$ in case 1 (birth after addition), as can be verified from (4). Let's examine the three remaining possibilities. Suppose step $i$ is removal of a $p$ simplex (cases 3 and 4). The removed simplex corresponds to a row of $\Phi_{p}^{i}$. Let $s_{i}$ be the indicator vector of that row. Then, by the description of the sequential algorithm in Section 3, to get $w_{i}^{T}$ we need to apply to $\Phi_{p}^{a ; \mathrm{v}}$ updates $a, \ldots, i-1$, select the row indicated by $s_{i}$ and columns of r, and negate the resulting submatrix. Formally (after some calculation which we skip), we get

$$
w_{i}^{T}=-s_{i}^{T} \Phi_{p}^{a ; \mathrm{v}}\left(\left[\begin{array}{c}
0 \\
I
\end{array}\right]+\left[\begin{array}{ccc}
u_{a}^{\prime} \cdots u_{i-1}^{\prime} & 0 \cdots 0 \\
0 \cdots 0 & 0 \cdots 0
\end{array}\right]\left[w_{a} \cdots w_{c}\right]^{T}\right)
$$

which implies

$$
\begin{aligned}
& \alpha_{i}^{T}=-s_{i}^{T} \Phi_{p}^{a ; v}\left[\begin{array}{c}
0 \\
I
\end{array}\right] \\
& \beta_{i}^{T}=-s_{i}^{T} \Phi_{p}^{a ; v}\left[\begin{array}{cccccc}
u_{a}^{\prime} & \cdots & u_{i-1}^{\prime} & 0 & \cdots & 0 \\
0 & \cdots & 0 & 0 & \cdots & 0
\end{array}\right] .
\end{aligned}
$$

Now suppose that step $i$ is addition of a $p$-simplex that kills a cycle (case 2). The killed cycle corresponds to a row in $\Gamma_{p}^{i}$. Let $s_{i}$ be the indicator vector of that row. Again, using the description of the sequential algorithm from Section 3 and skipping some calculations,

$$
\begin{aligned}
& w_{i}^{T}=-s_{i}^{T}\left(\left(\prod_{j=a}^{i-1} Q_{j}\right)^{T}+\right. \\
& \left.\left(\prod_{j=i}^{c} Q_{j}\right)\left[\begin{array}{llllll}
x_{a}^{\prime} & \cdots & x_{i-1}^{\prime} & 0 & \cdots & 0
\end{array}\right]\left[\begin{array}{llll}
y_{a}^{\prime} & \cdots & y_{c}^{\prime}
\end{array}\right]^{T}\right) \\
& \times \\
& \quad \Gamma_{p}^{a ; v}\left(\left[\begin{array}{l}
0 \\
I
\end{array}\right]+\left[\begin{array}{cccccc}
u_{a}^{\prime} & \cdots & u_{i-1}^{\prime} & 0 & \cdots & 0 \\
0 & \cdots & 0 & 0 & \cdots & 0
\end{array}\right]\left[\begin{array}{lll}
w_{a} & \cdots & w_{c}
\end{array}\right]^{T}\right)
\end{aligned}
$$

Letting

$$
\begin{aligned}
s_{i}^{\prime} & =\left(\prod_{j=a}^{i-1} Q_{j}\right) s_{i} \\
s_{i}^{\prime \prime} & =\left[\begin{array}{llllll}
x_{a}^{\prime} & \cdots & x_{i-1}^{\prime} & 0 & \cdots & 0
\end{array}\right]^{T}\left(\prod_{j=i}^{c} Q_{j}\right)^{T} s_{i} \\
s_{i}^{\prime \prime \prime} & =s_{i}^{\prime}+\left[\begin{array}{lll}
y_{a}^{\prime} & \cdots & y_{c}^{\prime}
\end{array}\right] s_{i}^{\prime \prime},
\end{aligned}
$$

we get

$$
\begin{aligned}
& \alpha_{i}^{T}=-\left(s_{i}^{\prime \prime \prime}\right)^{T} \Gamma_{p}^{a ; v}\left[\begin{array}{l}
0 \\
I
\end{array}\right], \\
& \beta_{i}^{T}=-\left(s_{i}^{\prime \prime \prime}\right)^{T} \Gamma_{p}^{a ; v}\left[\begin{array}{ccccccc}
u_{a}^{\prime} & \cdots & u_{i-1}^{\prime} & 0 & \cdots & 0 \\
0 & \cdots & 0 & 0 & \cdots & 0
\end{array}\right] .
\end{aligned}
$$

Appendix C gives pseudo-code for one possible efficient implementation of the above computations. Note that Left-to-Right also requires $\Phi_{p}^{a, \mathrm{v}}, \Gamma_{p}^{a, \mathrm{v}},\left\{\Phi_{p}^{i ; i}, \Gamma_{p}^{i ; i}: i \in \mathrm{l}\right\}$, in addition to already mentioned $\left\{M_{p}^{i ; 1}: i \in 1\right\}$. In the proof of correctness (Lemma 2) we will show that all this data is indeed available when Left-to-Right is invoked.

Batched application of updates. Next, we Apply-Left updates $M^{i ; \mathrm{v}}$ to matrices $\Phi^{a ; \mathrm{v}}$ and $\Gamma^{a ; \mathrm{v}}$ for all $i=a, \ldots, c$. The algorithm uses Prefix to compute $\left(\prod_{i=a}^{c} M_{p-1}^{i ; \mathrm{v}}\right)^{-1}$ and $\prod_{i=a}^{c} M_{p}^{i ; \mathrm{v}}$ as width- $k$ matrices (SUFFIX works just as well). Then it multiplies $\Phi_{p}^{a ; \mathrm{v}}$ and $\Gamma_{p}^{a ; \mathrm{v}}$ appropriately, resulting in $\Phi_{p}^{c+1 ; \mathrm{v}}, \Gamma_{p}^{c+1 ; \mathrm{v}}$.

## Apply-Left

for $p=0,1, \ldots$, max. dimension do
for $i \in l$ do
$P_{i}, u_{i}, v_{i} \leftarrow$ width- 1 form of $M_{p}^{i ; \mathrm{v}}$
$Q_{i}, x_{i}, y_{i} \leftarrow$ width- 1 form of $M_{p-1}^{i ; \mathrm{v}}$
$P, U^{\prime}, V^{\prime} \leftarrow \operatorname{Prefix}\left(P_{i}, u_{i}, v_{i} \mid i=a, \ldots, c\right)$
$Q, X^{\prime}, Y^{\prime} \leftarrow \operatorname{Prefix}\left(Q_{i}^{T}, \frac{Q_{i}^{T} x_{i}}{y_{i}^{T} Q_{i}^{T} x_{i}-1}, Q_{i} y_{i} \mid i=c, \ldots, a\right)$
$\Phi_{p}^{c+1 ; \mathrm{v}} \leftarrow \Phi_{p}^{a ; \mathrm{v}}\left(P+U^{\prime}\left(V^{\prime}\right)^{T}\right)$
$\Gamma_{p}^{c+1 ; \mathrm{v}} \leftarrow \Gamma_{p}^{a ; \mathrm{v}}\left(P+U^{\prime}\left(V^{\prime}\right)^{T}\right)$
$\Gamma_{p}^{c+1 ; \mathrm{v}} \leftarrow\left(Q+X^{\prime}\left(Y^{\prime}\right)^{T}\right) \Gamma_{p}^{c+1 ; \mathrm{v}}$
Right-to-Left. Right-to-Left extends the restriction of $\left\{M_{p}^{i}: i \in \mathrm{r}\right\}$ from r to v . Definition of restrictions of $M_{p}^{i}$ implies that the only rows/columns in $M_{p}^{i ; \mathrm{v}}$ which are not present in $M_{p}^{i ; \mathrm{r}}$ are those corresponding to $p$-cycles and $p$ chains removed in steps of 1 . In other words, $M_{p}^{i ; \mathrm{v}}$ is missing fewer topmost rows and leftmost columns compared to $M_{p}^{i, \mathrm{r}}$. But updates $\left\{M_{p}^{i}: i \in \mathrm{r}\right\}$ have no effect on $p$-cycles and $p$-chains removed in steps of $l$; this can be verified by inspecting update matrices in Section 3. Hence, we simply $\operatorname{pad}$ each $\left\{M^{i, \mathrm{r}}: i \in \mathrm{r}\right\}$, with an identity matrix of suitable size.

$$
\begin{aligned}
& \text { RIGHT-TO-LEFT } \\
& \text { for } i \in \mathrm{r} \text { and all } p \text { do } \\
& \qquad M_{p}^{i ; \mathrm{v}} \leftarrow\left[\begin{array}{cc}
I & 0 \\
0 & M_{p}^{i ; \mathrm{r}}
\end{array}\right]
\end{aligned}
$$

Apply-Right is implemented like Apply-Left, except $\left\{M_{p}^{i ; \mathrm{v}}: i \in \mathrm{l}\right\}$ are replaced by $\left\{M_{p}^{i ; \mathrm{v}}: i \in \mathrm{r}\right\}$, and $\Phi_{p}^{a, \mathrm{v}}, \Gamma_{p}^{a, \mathrm{v}}$ are replaced by $\Phi_{p}^{c+1 ; \mathrm{v}}, \Gamma_{p}^{c+1 ; \mathrm{v}}$.

Correctness. We prove correctness of the hierarchical algorithm by applying the following more general statement to the root of the operation tree.

Lemma 2. $\Phi^{a ; \mathrm{v}}, \Gamma^{a ; \mathrm{v}}$ are correct when $\mathrm{H}-\mathrm{ZZPH}(\mathrm{v})$ is invoked and thereafter. $\left\{M^{i, \mathrm{v}}, i \in \mathrm{v}\right\}, \Phi^{b+1 ; \mathrm{v}}, \Gamma^{b+1 ; \mathrm{v}}$ are correct when $\mathrm{H}-\mathrm{ZZPH}(\mathrm{v})$ terminates and thereafter.

Proof. We prove the statement by induction on the order of events (invocations and terminations of H-ZZPH).

The first event is invocation where v is the root. In that case, $\Phi_{p}^{a ; \mathrm{v}}$ and $\Gamma_{p}^{a ; \mathrm{v}}$ are simply initial unrestricted matrices $\Phi_{p}^{1}$ and $\Gamma_{p}^{1}$, which are correct initially, because they are initialized in the same way as in the sequential algorithm.
Now consider some other event - an invocation or a termination at a node $u$ - assuming that the claim is true for all the preceding events.

Suppose that the event is an invocation of $\mathrm{H}-\mathrm{ZZPH}(u)$. Noting that $u$ is not a root, let v be the parent of $u$. If $u=1$, since the restrictions of the matrices to a child are a subset of the restrictions to a parent, we are done by the inductive hypothesis (i.h.) on the invocation of $\mathrm{H}-\mathrm{ZZPH}(\mathrm{v})$. Otherwise, by i.h., when H-ZZPH(1) terminates, $\left\{M^{i ; 1}: i \in\right.$ $1\}, \Phi^{c+1 ; 1}, \Gamma^{c+1 ; 1}$ are correct. By i.h. for the invocation on each leaf $i \in \mathrm{l},\left\{\Phi_{p}^{i ; i}, \Gamma_{p}^{i ; i}: i \in \mathrm{l}\right\}$ are correct. Hence the vectors $s_{i}$ in Left-TO-Right can be correctly read off from $\Phi_{p}^{i ; i}$ and $\Gamma_{p}^{i ; i}$. Since all input data for Left-to-Right is correct, $\left\{M^{i, \mathrm{v}}: i \in \mathrm{l}\right\}$ are correct when Left-To-Right terminates. By design of Apply-Left, at its termination, $\Phi^{c+1, \mathrm{v}}, \Gamma^{c+1 ; \mathrm{v}}$, and, therefore, $\Phi^{c+1, \mathrm{r}}, \Gamma^{c+1 ; \mathrm{r}}$ are correct, which completes this part of the proof.

Now suppose that the event is termination of $\mathrm{H}-\mathrm{ZZPH}(\mathrm{v})$. If v is a leaf, then at termination, $M^{a ; \mathrm{v}}, \Phi^{a+1 ; \mathrm{v}}, \Gamma^{a+1 ; \mathrm{v}}$ are correct because they are computed using the same rules as in the sequential algorithm, and starting from correct input ( $\Phi^{a ; \mathrm{v}}, \Gamma^{a ; \mathrm{v}}$ ). Now suppose v is internal. As before, combining i.h. on termination of $\mathrm{H}-\mathrm{ZZPH}(\mathrm{l})$, i.h. on the invocation on the leaf descendents of l, and design of Left-to-Right and Apply-Left, we prove that when ApplyLeft terminates, $\left\{M^{i ; \mathrm{v}}: i \in \mathrm{l}\right\}, \Phi^{c+1 ; \mathrm{v}}, \Gamma^{c+1 ; \mathrm{v}}$ are correct. By i.h., when H-ZZPH(r) terminates, $\left\{M^{i ; \mathrm{r}}: i \in \mathrm{r}\right\}, \Phi^{b+1 ; \mathrm{r}}$, $\Gamma^{b+1 ; \mathrm{r}}$ are correct. When H-ZZPH(v) terminates, $\left\{M^{i ; \mathrm{v}}\right.$ : $i \in \mathrm{r}\}$ are correct by design of Right-To-Left, and $\Phi^{b+1 ; \mathrm{v}}$, $\Gamma^{b+1 ; \mathrm{v}}$ are correct by design of Apply-Right.

Running time. We show that the hierarchical algorithm runs on an $n$-step zigzag in time $O\left(M(n)+n^{2} \log ^{2} n\right)$. The key step is showing that $\mathrm{H}-\mathrm{ZZPH}(\mathrm{v})$, excluding its recursive calls, can be implemented to run in time proportional to the number of leaf descendents of v . The missing proofs of lemmas below appear in Appendix B.

Theorem 1. H-ZZPH(v) runs in $O\left(\frac{n}{k} M(k)+\frac{n}{k} k^{2} \log ^{2} k\right)$ time.

From this we easily get the main result of the paper.
Corollary 1. Zigzag persistent homology for a sequence of $n$ steps can be computed in time $O\left(M(n)+n^{2} \log ^{2} n\right)$.

Before proving Theorem 1, we provide tools for efficient computation that will be invoked repeatedly.

Lemma 3. Let $U$, $V$ be $n \times k$ matrices, $k \leq n$, and let $W$ be $a$ $k \times k$ matrix. One can compute $U^{T} V$ and $U W$ in $O\left(\frac{n}{k} M(k)\right)$ time.

Lemma 4. Let $B$ be an $n \times k$ matrix. Given a width- $k$ representation of $A, A B$ can be computed in $O\left(\frac{n}{k} M(k)\right)$ time.

Lemma 5. If $A$ has width $k_{1}$ and $B$ has width $k_{2}$, then $A B$ has width $k_{1}+k_{2}$. Given low-width representations $A$ and $B$, low-width representation of $A B$ can be computed in $O\left(\frac{n}{k_{1}+k_{2}} M\left(k_{1}+k_{2}\right)\right)$ time.

Proof of Theorem 1. First assume that $M(n) \notin O\left(n^{2}\right)$.
Let $T_{\text {Prefix }}(t)$ be the worst-case running time of Prefix on an input consisting of $t$ width- 1 terms. Then $T_{\text {PRefix }}(t) \leq$ $2 T_{\text {Prefix }}\left(\frac{t}{2}\right)+O\left(\frac{n}{t} M(t)\right)$, where the last term is determined by the time it takes to compute $Y_{1}^{T} X_{2}$ by Lemma 3 . Solving this recursion, we get $T_{\text {Prefix }}(t)=O\left(\frac{n}{t} M(t)\right)$. By similar
analysis we can bound the running time of Suffix on $t$ terms by $O\left(\frac{n}{t} M(t)\right)$.

Apply-Left and Apply-Right run in time $2 T_{\text {Prefix }}(k)+$ $O\left(\frac{n}{k} M(k)\right)+O\left(\frac{n}{2 k} M(2 k)\right)+O\left(\frac{n}{3 k} M(3 k)\right)$, where the terms correspond to two calls to Prefix, computing $\Phi_{p}^{a ; \mathrm{v}}(P+$ $\left.U^{\prime}\left(V^{\prime}\right)^{T}\right)$ by Lemma 4 , computing $\Gamma_{p}^{a ; \mathrm{v}}\left(P+U^{\prime}\left(V^{\prime}\right)^{T}\right)$ by Lemma 5 , and computing $\left(Q+X^{\prime}\left(Y^{\prime}\right)^{T}\right) \Gamma_{p}^{c+1 ; \mathrm{v}}$ by Lemma 5 .

We analyze the running time of Left-to-Right, as implemented in Appendix C. Let $k_{p}$ be the number of nontrivial updates for dimension $p$. Clearly, $\sum_{p} k_{p}=O(k)$. We fix dimension $p$, and analyze its iteration of the for loop. Calls to Prefix and Suffix take $O\left(\frac{n}{k} M(k)\right)$ time. Computing $s_{i}^{\prime}$ and $s_{i}^{\prime \prime}, i \in 1$ takes $O(n k)$ time, using any obvious implementation. Computing $Y^{\prime}\left[\begin{array}{lll}s_{a}^{\prime \prime} & \cdots & s_{c}^{\prime \prime}\end{array}\right]$ takes $O\left(\frac{n}{k} M(k)\right)$ time by Lemma 3. Computing [ $\left.\begin{array}{ccc}s_{a}^{\prime \prime \prime} & \cdots & s_{c}^{\prime \prime \prime}\end{array}\right]$ takes $O(n k)$ time. Since $\Phi_{p}^{a ; \mathrm{v}}$ has size at most $k \times n$, computing $\Phi_{p}^{a ; \mathrm{v}} U^{\prime}$ takes $O\left(\frac{n}{k} M(k)\right)$ by Lemma 3. Since $\Gamma_{p}^{a ; \mathrm{v}}$ has width $k$, computing $\Gamma_{p}^{a ; \mathrm{v}} U^{\prime}$ takes $O\left(\frac{n}{k} M(k)\right)$ time by Lemma 4. Multiplying the result by $\left[\begin{array}{llll}s_{1}^{\prime \prime \prime} & \cdots & s_{k}^{\prime \prime \prime}\end{array}\right]^{T}$ takes $O\left(\frac{n}{k} M(k)\right)$ time, by Lemma 3. Coefficient vectors $\alpha_{i}^{T}, \beta_{i}^{T}$ are computed in $O\left(k^{2}\right)$ time, because they appear as submatrices of already computed matrices $T_{1}$ and $T_{2}$. The $k \times k$ linear system can be solved in $O(M(k))$ time using Bunch and Hopcroft's algorithm [11]. We conclude that the for loop for dimension $p$ runs in $O\left(\frac{n}{k} M(k)\right)$ time. It can actually be implemented to run in $O\left(\frac{n}{k_{p}} M\left(k_{p}\right)\right)$, where $k_{p}$ is the number of updates that affect dimension $p$. Since this is superlinear in $k_{p}$, and $\sum_{p} k_{p}=O(k)$, we conclude that Left-to-Right runs in $O\left(\frac{n}{k} M(k)\right)$ time.

Right-To-Left modifies $k$ matrices by changing adding $k$ nonzero entries to each. Clearly, it can be implemented to run in optimal $O\left(k^{2}\right)$.

We define $T_{\mathrm{H}-\mathrm{ZZPH}}(k)$ to be the worst-case running time of $\mathrm{H}-\mathrm{ZZPH}(\mathrm{v})$ for any tree node v with $k$ leaf descendents. It is easy to see that $T(1)=O(n)$, because if v is a leaf, the algorithm manipulates $O(1)$ rows and columns, by inspecting, moving or copying them to compute $M_{p}^{a ; a}, \Phi_{p}^{a+1 ; a}$, $\Gamma_{p}^{a+1 ; a}$. Furthermore, from the above analysis it follows that $T_{\mathrm{H}-\mathrm{ZZPH}}(2 k) \leq 2 T_{\mathrm{H}-\mathrm{ZZPH}}(k)+O\left(\frac{n}{k} M(k)\right)$. The claim follows by solving the recurrence relation.

If $M(n) \in O\left(n^{2}\right)$ the analysis is similar. Solving each recurrent relation (the one for $T_{\text {Prefix }} / T_{\text {SUffix }}$, and the one for $T_{\mathrm{H}-\mathrm{ZZPH}}$ ) contributes one logarithmic factor.

## Acknowledgement

The authors would like to thank Leonidas Guibas, Gunnar Carlsson and Frédéric Chazal for their support. This work has been supported in part by NSF grants CCF 0634803, FODAVA 0808515, CCF 1011228, DARPA grant HR0011-05-1-0007 and ANR-09-BLAN-0331-01.

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## APPENDIX

## A. CORRECTNESS OF THE SEQUENTIAL ALGORITHM

Since the operations in Section 3 are almost the same as in [6], we focus our proof of correctness on verifying the new parts of the invariant.

1: $D \Phi_{p}^{j}=\Phi_{p-1}^{j} \Gamma_{p}^{j}$. The equality is satisfied in Equation (3) because we apply the same updates to both sides of the equation. However, we must still ensure that the specific forms prescribed to matrices $\Phi_{p}^{j}$ and $\Gamma_{p}^{j}$ remain intact: namely, that the blocks marked zero in (3) remain zero.

If there is a birth after an addition, the product $\Phi_{p}^{j} M_{p}^{j}$ moves the newly born cycle from submatrix $F_{p}^{j}$ into $Z_{p}^{j+1}$; its boundary is zero by the case assumption. $\Gamma_{p}^{j} M_{p}^{j}$ moves the zero column (zero by Lemma 1) into the portion of $\Gamma_{p}^{j+1}$ corresponding to the cycle matrix $Z_{p}^{j}$. Pre-multiplying $\Gamma_{p+1}$ by $\left(M_{p}^{j}\right)^{-1}$ performs the corresponding row update, safely moving a row from the fourth to the second section of $\Gamma_{p+1}$. If there is a death after an addition, $M_{p-1}^{j}$ performs a basis change in $Z_{p-1}^{j}$ (with a matching update in $B_{p}^{j}$ ). It does not affect the structure of the matrices, only ensures that the leftmost column of $B_{p}^{j+1}$ has a single 1. $M_{p}^{j}$ does two things. First it moves the bounding chain from $F_{p}^{j}$ into $C_{p}^{j+1}$, and the corresponding boundary from $H_{p}^{j}$ to $B_{p}^{j+1}$. Then it subtracts a multiple of the new column in $B_{p}^{j+1}$ from $H_{p}^{j}$; these subtractions only introduce zeros. The effect of $\left(M_{p}^{j}\right)^{-1}$ on $\Gamma_{p+1}^{j}$ is adding rows of $L_{p+1}^{j}$ to $K_{p+1}^{j}$, a benign update as far as zeros are concerned.
If a simplex is removed, multiplying by matrix $M_{p}^{j}$ zeroes out the outgoing row of matrix $C_{p}^{j}$ (and also $Z_{p}^{j}$ in case of a death), as well as $F_{p}^{j}$. As a result, once the row is moved to the past, all but its first portion (now belonging to matrix $E_{p}^{j+1}$ ) is zero. The effect of the matrix $\left(M_{p-1}^{j}\right)^{-1}$ (from Equation (6)) in the product $\left(M_{p-1}^{j}\right)^{-1} \Gamma_{p}^{j}$ does not cross the partition boundaries.

2: $B_{p}^{j}$ has exactly one non-zero element per column, and at
most one per row. $H_{p}^{j}$ is independent of $B_{p}^{j}$. In the removal cases, the only change to the matrix $H_{p}^{j}$ is that it loses a row, and so does $B_{p}^{j}$; this loss does not affect the invariant. It's important to note that in case of the change of basis in $Z_{p}^{j}$ after a death, $B_{p}^{j}$ cannot stop satisfying the invariant, since the columns of $Z_{p}^{j}$ that have a matching 1 in $B_{p}^{j}$ cannot contain the outgoing simplex (the 1 would indicate that the cycles were boundaries, but the outgoing simplex has no cofaces). In case of a birth after removal, $M_{p-1}^{j}$ performs a change of basis in $Z_{p-1}^{j+1}$ such that its inverse updates $B_{p}^{j+1}$ to once again satisfy the invariant.

In the case of addition, if birth occurs, matrix $H_{p}^{j}$ loses a zero column, which again requires no update. If there is a death, $B_{p}^{j}$ acquires a single column. Pre-multiplication of $\Gamma_{p}^{j}$ by $\left(M_{p-1}^{j}\right)^{-1}$ ensures that $B_{p}^{j+1}$ has a single 1 in that column, which is in the row of the lowest non-zero entry in the outgoing column of $H_{p}^{j}$; therefore, $B_{p}^{j+1}$ remains a permutation matrix. We explicitly zero this row out via matrix $M_{p}^{j}$. Therefore, in all cases statement 2 of the Invariant is maintained.

3, 4: Matrix $M_{p}^{j}$ perform performs additions from left to right in $F_{p}^{j}$, and possibly removes a column from it (in the case of an addition of a simplex). Therefore, the columns of $F_{p}^{j+1}$ remain linearly independent.

In the case of a removal of a simplex, the main updates performed to matrices $Z$ and $C$ are the same as in [6], with the exception of a few additional changes of basis in $Z$ via left to right operations (which respect the right filtration). Therefore, the last two parts of the invariant are maintained.

In the case of an addition, the only difference from [6] is that the first column of the matrix $F_{p}^{j}$ already has the correct form, and does not require any additional processing.

In the birth case, this is easy to see: the column $F_{p}^{0}[i]$ corresponding to the simplex $\sigma_{i}$ contains a 1 in the row corresponding to this simplex. None of the operations ever remove this element from this column. Therefore, when we add $\sigma_{i}$ we have a new cycle (recorded as the column $F_{p}^{j}[i]$ ), which we append to the matrix $Z_{p}^{j}$.

In the death case, the boundary of the column $F_{p}^{j}[i]$ is in the kernel of the map on the homology. If the lowest element in the matching column $H_{p}^{j}[i]$ is in the row $l$, then this cycle is in the span of the first $l$ elements of the right filtration given by the matrix $Z_{p-1}^{j}$. Furthermore, since the matrix $H_{p}^{j}$ is reduced with respect to $B_{p}^{j}$, the row $l$ in $B_{p}^{j}$ is zero, by definition. Therefore, there does not exist a cycle in the span of the first $l-1$ elements of the right filtration that's in the kernel of the homology map. Therefore, our update is correct - it adds to the matrix $B_{p}^{j}$ a column with a 1 in row $l$, indicating that the updated cycle $Z_{p-1}^{j+1}$ dies after the addition of simplex $\sigma_{i}$. (This analysis reflects the Reduction Lemma in [6].) In summary, the operations in Section 3 closely mimic the original zigzag persistent homology algorithm [6], and satisfy the additional Sequential Invariant requirements.

## B. PROOFS OF RUNNING TIME LEMMAS

Proof of Corollary 1. This is a special case for $k=n$ (i.e. root of the tree), with $Z_{p}, C_{p}, B_{p}, H_{p}$ and $K_{p}$ "empty" ( $Z_{p}$, $C_{p}, B_{p}$ with zero columns, $H_{p}, K_{p}$ with zero rows), with $F_{p}$ equal to a permutation matrix that maps addition order to removal order, and $L_{p}=F_{p-1}^{T} D_{p} F_{p}$.

Proof sketch of Lemma 3. Split matrices into $k \times k$ blocks, and treat blocks as entries. Use a fast $k \times k$ matrix multiplication algorithm to multiply blocks.

Proof sketch of Lemma 5. Let $P, U, V$, and $Q, X, Y$ be low-width representations of $A$ and $B$, respectively. It is easy to check that

$$
P Q, \quad\left[\begin{array}{ll}
U & P X+U\left(V^{T} X\right)
\end{array}\right], \quad\left[\begin{array}{cc}
Q^{T} V & Y
\end{array}\right]
$$

is a width- $\left(k_{1}+k_{2}\right)$ representation of $A B$. The running time follows by applying Lemma 3 to the above expressions.

Proof sketch of Lemma 4. Let $P, U, V$ be the width- $k$ representation of $A$. Then one can compute $A B$ as $P B+$ $U\left(V^{T} B\right)$, using Lemma 3 .

## C. LEFT-TO-RIGHT IMPLEMENTATION

The following is one possible implementation of the Left-to-Right computation, described in Section 4, expressed in pseudo-code.

$$
\begin{aligned}
& \text { Left-To-Right(v) } \\
& \text { for } i \in l \text { do } \\
& Q_{i} \leftarrow I, \quad x_{i} \leftarrow 0, \quad y_{i} \leftarrow 0 \\
& \text { for } p=0,1, \ldots \text {, maximum dimension do } \\
& \text { for } i=a, \ldots, c \text { do } \\
& P_{i}, u_{i}, v_{i} \leftarrow \text { width- } 1 \text { form of } M_{p}^{i, 1} \\
& s_{i} \leftarrow \text { pivot row of } M_{p}^{i, 1} \text { (if any) } \\
& P, U^{\prime}, V^{\prime} \leftarrow \operatorname{Prefix}\left(P_{i}, u_{i}, v_{i} \mid i=a, \ldots, c\right) \\
& Q, X^{\prime}, Y^{\prime} \leftarrow \operatorname{SuFFIX}\left(Q_{i}^{T}, \frac{Q_{i}^{T} x_{i}}{y_{i}^{T} Q_{i}^{T} x_{i}-1}, Q_{i} y_{i} \mid i=c, \ldots, a\right) \\
& T_{1} \leftarrow I, \quad T_{2} \leftarrow Q \\
& \text { for } i=a, \ldots, c \text { do } \\
& s_{i}^{\prime} \leftarrow T_{1} s_{i} \\
& s_{i}^{\prime \prime} \leftarrow\left[\begin{array}{ll}
e_{1} \cdots e_{i-a} & 0 \cdots 0
\end{array}\right]^{T}\left(X^{\prime}\right)^{T}\left(T_{2} s_{i}\right) \\
& T_{1} \leftarrow T_{1} Q_{i}, \quad T_{2} \leftarrow T_{2} Q_{i} \\
& T_{1} \leftarrow Y^{\prime}\left[\begin{array}{lll}
s_{a}^{\prime \prime} & \cdots & s_{c}^{\prime \prime}
\end{array}\right] \\
& {\left[\begin{array}{ccc}
s_{a}^{\prime \prime \prime} & \ldots & s_{c}^{\prime \prime \prime} \\
T_{1} & \Phi^{a ;{ }^{a}} U^{\prime}
\end{array}\right] \leftarrow\left[\begin{array}{ccc}
s_{a}^{\prime} & \cdots & s_{c}^{\prime} \\
T_{0} & \leftarrow \Gamma^{a ; \mathrm{v}} U^{\prime}
\end{array}\right]+T_{1}}
\end{aligned}
$$

$$
\begin{aligned}
& T_{2} \leftarrow\left[\begin{array}{lll}
s_{a}^{\prime \prime \prime} & \cdots & s_{c}^{\prime \prime \prime}
\end{array}\right]^{T} T_{2} \\
& \text { for } i \in 1 \text { do } \\
& {\left[\alpha_{i}^{T} \mid \beta_{i}^{T}\right] \leftarrow 0 \text { or } s_{i}^{T} T_{1} \text { or } e_{i-a+1}^{T} T_{2}} \\
& \beta_{i}^{T} \leftarrow \beta_{i}^{T}\left[\begin{array}{ll}
e_{1} \cdots e_{i-a} & 0 \cdots 0
\end{array}\right] \\
& {\left[w_{a} \cdots w_{c}\right] \leftarrow-\left[\beta_{a} \cdots \beta_{c}\right]\left(\left[\alpha_{a} \cdots \alpha_{c}\right]-I\right)^{-1}} \\
& \text { for } i=a, \ldots, c \text { do } \\
& Q_{i} \leftarrow\left[\begin{array}{cc}
P_{i} & 0 \\
0 & I
\end{array}\right] \\
& x_{i} \leftarrow\left[\begin{array}{c}
u_{i} \\
0
\end{array}\right], \quad y_{i} \leftarrow\left[\begin{array}{c}
v_{i} \\
w_{i}
\end{array}\right] \\
& M_{p}^{i ; \mathrm{v}} \leftarrow Q_{i}+x_{i} y_{i}^{T}
\end{aligned}
$$


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    SCG'11, June 13-15, 2011, Paris, France.
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[^1]:    H-ZZPH(v)
    if v is leaf then
    Compute $M_{p}^{a ; a}, \Phi_{p}^{a+1 ; a}, \Gamma_{p}^{a+1 ; a}$ as in Section 3, but restricting to suitable $1 \times 1,1 \times n$, or $n \times 1$ submatrices
    if v is internal then H-ZZPH(1) Left-to-Right Apply-Left H-ZZPH(r) Right-to-Left Apply-Right

