

Computation of persistent homology

Ulrich Bauer

August 13, 2018

1 Computing homology

We will now describe a matrix algorithm for computing (persistent) homology with coefficients in a fixed field \mathbb{F} . As usual, we assume $\mathbb{F} = \mathbb{Z}_2$ for simplicity, mentioning however when other choices of coefficients require modifications. The basic idea is to find a compatible basis for the subspaces $B_*(K) \subseteq Z_*(K) \subseteq C_*(K)$.

Remark 1. Let $\Sigma_U \subset \Sigma_V$ be bases of vector spaces $U \subset V$. Then $(\Sigma_V \setminus \Sigma_U)$ is a set of representatives for a basis of the quotient space V/U .

Proof. Let $\Sigma_U = \{v_1, \dots, v_m\}$, $\Sigma_V = \{v_1, \dots, v_n\}$. We first show that $\Sigma_W = \{v_{m+1} + U, \dots, v_n + U\}$ spans V/U , and then show that Σ_W is linearly independent.

Let $v \in V$. Then there are unique coefficients λ_i such that $v = \sum_{i=1}^n \lambda_i v_i$. Now

$$v + U = \sum_{i=1}^n \lambda_i v_i + U = \sum_{i=m+1}^n \lambda_i (v_i + U),$$

and thus Σ_W spans V/U .

In order to show that Σ_W are linearly independent, we need to show that $\lambda_i = 0$ for $i = m + 1, \dots, n$ whenever $v + U = 0$, i.e., whenever

$$\sum_{i=m+1}^n \lambda_i v_i \in U.$$

Since Σ_U is a basis for U , we get $(\lambda_i)_{i=1}^m$ such that

$$\sum_{i=1}^m \lambda_i v_i = \sum_{i=m+1}^n \lambda_i v_i.$$

But the v_i are linearly independent, and thus $\lambda_i = 0$ for all $i = 1, \dots, n$. \square

Consider a simplicial complex K . The *boundary matrix* of K is the matrix D of the boundary map $\partial : C_*(K) \rightarrow C_*(K)$ with respect to the basis given by the simplices $\{\sigma_1, \dots, \sigma_n\}$ of K .

Definition 2. Let M be a matrix. The *pivot index* of a column m_i of M is

$$\text{pivot}(m_i) = \min\{j \in \mathbb{N} : m_{ik} = 0 \text{ for all } k > j\}.$$

We write

$$\text{pivots } M = \{\text{pivot } m_j : 1 \leq j \leq n\} \setminus \{0\}.$$

A matrix M is *reduced* if the pivots of non-zero columns are distinct, i.e., $i \neq j$ implies either $\text{pivot}(m_i) \neq \text{pivot}(m_j)$ or $0 = \text{pivot}(m_i) = \text{pivot}(m_j)$.

Lemma 3. *The nonzero columns of a reduced matrix M are linearly independent.*

Proof. Let $(\lambda_j)_j$ be such that

$$v = \sum_j \lambda_j m_j = 0.$$

Then

$$0 = \text{pivot } v = \max\{\text{pivot } m_j \mid \lambda_j \neq 0\}$$

by the assumption that the pivots are unique. Thus, $m_j \neq 0$ implies $\lambda_j = 0$, meaning that the nonzero columns are linearly independent. \square

Homology with field coefficients can be computed using a variant of Gaussian elimination. Specifically, the following algorithm finds an upper triangular matrix V such that $R = D \cdot V$ is reduced.

Algorithm 4 (matrix reduction).

Require: D : $m \times n$ matrix

Ensure: $R = D \cdot V$ is reduced, V is full rank upper triangular

function REDUCE(D)

$R = D, V = I(n)$

while there exist $i < j$ with $\text{pivot } R_i = \text{pivot } R_j$ **do**

 add column R_i to column R_j

 add column V_i to column V_j

end while

return R, V

end function

Note that the reduced matrix $R = D \cdot V$ is not unique, and the algorithm is not deterministic. To obtain a deterministic variant of the algorithm, one may iterate over $j = 1, \dots, n$. For coefficients other than \mathbb{Z}_2 , the columns R_i and V_i have to be rescaled so that their pivot entries cancel to 0 in the column addition.

Example 5. Consider the empty triangle

$$K = \{\{a\}, \{b\}, \{c\}, \{a, b\}, \{a, c\}, \{b, c\}\}.$$

The boundary matrix is

$$D = \begin{pmatrix} 0 & 0 & 0 & 1 & 1 & 0 \\ 0 & 0 & 0 & 1 & 0 & 1 \\ 0 & 0 & 0 & 0 & 1 & 1 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \end{pmatrix}$$

and a matrix reduction is given by

$$R = \begin{pmatrix} 0 & 0 & 0 & 1 & 1 & 0 \\ 0 & 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \end{pmatrix} = D \cdot \begin{pmatrix} 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 & 0 & 1 \\ 0 & 0 & 0 & 0 & 1 & 1 \\ 0 & 0 & 0 & 0 & 0 & 1 \end{pmatrix}.$$

We have pivots $R = \{2, 3\}$; the non-zero column indices of R are $N = \{4, 5\}$, the zero column indices are $P = \{1, 2, 3, 6\}$, and the non-pivot zero column indices are $E = P \setminus \text{pivots } R = \{1, 6\}$.

Theorem 6. *The homology of K can be computed by reducing the boundary matrix D .*

(a) *The columns*

$$\Sigma_B = \{r_j \neq 0\}$$

form a basis of $B_(K)$.*

(b) The columns

$$\widetilde{\Sigma}_Z = \{V_i \mid r_i = 0\}$$

form a basis of $Z_*(K)$.

(c) Exchanging each basis cycle

$$v_i \in \widetilde{\Sigma}_Z \text{ with } i \in \text{pivots } R$$

with the basis boundary

$$r_j \in \Sigma_B \text{ with } i = \text{pivot } r_j = \text{pivot } v_i$$

extends Σ_B to another basis of $Z_*(K)$,

$$\Sigma_Z = \Sigma_B \cup \Sigma_E$$

with

$$\Sigma_E = \{v_i \mid r_i = 0, i \notin \text{pivots } R\} \subseteq \widetilde{\Sigma}_Z.$$

(d) The non-bounding basis cycles Σ_E generate a basis

$$\Sigma_H = \{[v_i] = v_i + B_d(K) \mid r_i = 0, i \notin \text{pivots } R, d = \dim \sigma_i\}$$

for $H_*(K)$.

(e) The Betti numbers $\beta_d(K)$ can be read off from R , without knowing V :

$$\beta_d = \text{card}\{i \in E : \dim \sigma_i = d\}$$

(f) We can further extend Σ_Z to a basis for $C_*(K)$:

$$\Sigma_C = \Sigma_Z \cup \{v_j \mid r_j \neq 0\}$$

$$= \{r_j \mid r_j \neq 0\} \cup \{v_j \mid r_j \neq 0\} \cup \{v_i \mid r_i = 0, i \notin \text{pivots } R\}.$$

The matrix of ∂ with respect to this basis has at most one nonzero entry in each row/column.

Proof. (a) The columns of the boundary matrix D are a generating set for B_* by definition. Since V is regular, the columns of $R = D \cdot V$ are a generating set as well. The columns in Σ_B have distinct pivot indices and therefore have to be linearly independent, by Lemma 3. Therefore the columns Σ_B form a basis.

(b) Let $z \in Z_*$. Since V is regular, its columns $\{v_i\}_i$ form a basis for $C_*(K)$, and we can write

$$z = \sum_i \lambda_i v_i.$$

Now

$$0 = \partial z = \sum_i \lambda_i \partial v_i = \sum_i \lambda_i r_i$$

implies, with the fact that Σ_B is a basis, that

$$\lambda_j = 0$$

for all j with $r_j \neq 0$. Therefore we can write

$$z = \sum_{i:r_i=0} \lambda_i v_i,$$

and so $\widetilde{\Sigma}_Z$ is a generating set for $Z_*(K)$. The columns v_i have again distinct pivot indices and are thus linearly independent. We conclude that $\widetilde{\Sigma}_Z$ is a basis for $Z_*(K)$.

(c) Replacing all v_i for $i = \text{pivot } r_j = \text{pivot } v_i$ yields pivots $\widetilde{\Sigma}_Z = \text{pivots } \Sigma_Z$, so Σ_Z also has distinct pivots and is therefore also a maximal linearly independent set of cycles. Thus Σ_Z is another basis for $Z_*(K)$.

(d) This follows from Remark 1 applied to $\Sigma_B \subseteq \Sigma_Z$.

- (e) The dimension $\beta_d(K)$ of $H_d(K)$ is the cardinality of its basis.
- (f) Similarly to before, Σ_C is obtained from the columns of V by exchanging vectors with the same pivot index, and therefore Σ_C has distinct pivots and is thus a basis. We have $\partial v_j = r_j$ for all $r_j \neq 0$, while all other basis elements are cycles, implying the claim. \square

2 Persistent Homology

As it turns out, the same algorithm can also be used to compute the persistent homology of a filtration.

Consider a *simplexwise filtration* K_\bullet of a simplicial complex K such that $K_i = K_{i-1} \cup \{\sigma_i\}$ for $1 \leq i \leq n$, where $\sigma_i \in K$ is a simplex. The *boundary matrix* of K_\bullet is the matrix D of the boundary map $\partial : C_*(K) \rightarrow C_*(K)$ with respect to the ordered basis $(\sigma_i)_{1 \leq i \leq n}$. In particular, D is the boundary matrix of the final complex K .

Remark 7. The persistent homology of K_\bullet can be computed by reducing the boundary matrix D :

- (a) The columns

$$\Sigma_{B,k} = \{r_j \neq 0 \mid j \leq k\}$$

form a basis of $B_*(K_k)$.

- (b) The columns

$$\widetilde{\Sigma}_{Z,k} = \{V_i \mid r_i = 0, i \leq k\}$$

form a basis of $Z_*(K)$.

- (c) Exchanging each basis cycle

$$v_i \in \widetilde{\Sigma}_Z \text{ with } i \leq k \in \text{pivots } R$$

with the basis boundary

$$r_j \in \Sigma_B \text{ with } i = \text{pivot } r_j$$

extends $\Sigma_{B,k}$ to another basis of $Z_*(K_k)$,

$$\Sigma_{Z_k} = \Sigma_{B_k} \cup \Sigma_{P_k} \cup \Sigma_{E_k}$$

with

$$\begin{aligned} \Sigma_{P,k} &= \{r_j \neq 0 \mid i = \text{pivot } r_j, k \in [i, j]\}, \\ \Sigma_{E,k} &= \{v_i \mid r_i = 0, i \neq \text{pivots } R, k \in [i, \infty)\}. \end{aligned}$$

(d) The additional cycles generate a basis

$$\Sigma_{H,k} = \{[z] \mid z \in \Sigma_{P,k} \cup \Sigma_{E,k}\}$$

for the homology $H_*(K_k)$.

(e) The inclusion-induced map $H_*(K_k) \rightarrow H_*(K_l)$ has rank

$$\text{card}(\Sigma_{H_k} \cap \Sigma_{H_l}).$$

Remark 8. Persistent homology (of a simplexwise filtration indexed by natural numbers $[n] = \{1, \dots, n\}$) is a (commutative) diagram of vector spaces $V_i = H_*(K_i)$, indexed by $[n]$, and linear maps $f_{ij} : V_i \rightarrow V_j$ ($i \leq j$) with $f_{ik} = f_{jk} \circ f_{ij}$ ($i \leq j \leq k$) and $f_{ii} = \text{Id}_{V_i}$. Such a diagram is also called a *persistence module*.

Definition 9. The collection of intervals $B = B(H_*(K_\bullet))$,

$$B = \{[i, \infty) \mid r_i = 0, i \notin \text{pivots } R\} \cup \{[i, j) \mid i = \text{pivot } r_j\}$$

is the *persistence barcode* of the filtration K_\bullet .

Corollary 10. Let \mathbb{F} be a field. For any interval $I = [i, j) \subset \mathbb{N}$ or $I = [i, \infty) \subset \mathbb{N}$, define a diagram $\mathbf{R} \rightarrow \mathbf{Vect}$,

$$\mathbb{F}_k^I = \begin{cases} \mathbb{F} & \text{if } k \in I, \\ 0 & \text{otherwise,} \end{cases}$$

with all maps $\mathbb{F}_k^I \rightarrow \mathbb{F}_l^I$ ($k \leq l$) having maximal rank. Then

$$H_*(K_\bullet) \cong \bigoplus_{I \in B(H_*(K_\bullet))} \mathbb{F}_\bullet^I.$$

3 Reindexing

While we have seen examples of general filtrations indexed over \mathbb{R} , our algorithm requires a simplexwise filtration indexed over $[n] = \{1, \dots, n\}$. We will now describe a straightforward way of applying this algorithm to the more general setting.

There are two separate issues: transforming the indexing from \mathbb{R} to $[n]$, and transforming a general filtration to a simplex-wise one.

Definition 11. We say that a diagram $G : \mathbb{R} \rightarrow C$ (in any category C) is *essentially finite* if there exists a diagram $F : [n] = \{1, \dots, n\} \rightarrow C$ and a monotonic reindexing map $r : \mathbb{R} \rightarrow [n]$ such that $G = F \circ r$.

This way we can interpret G as a diagram H indexed over $[n]$, letting $H_i = G_i$ for any $t \in T_i$. We can also extend the index category to \mathbb{N} by setting $H_m = H_n$ for $m > n$, connected by the identity. Conversely, given the \mathbf{n} -indexed diagram together with the interval partition, we can recover the \mathbf{R} -indexed diagram.

Definition 12. We say that a simplexwise filtration $L_\bullet : \mathbb{N} \rightarrow \mathbf{Simp}$ *refines* a filtration $K_\bullet : \mathbb{N} \rightarrow \mathbf{Simp}$ if there is a map $m : \mathbb{N} \rightarrow \mathbb{N}$ such that $K_i = L_{m(i)}$ for all $i \in \mathbb{N}$.

Given a filtration $K_\bullet : \mathbb{N} \rightarrow \mathbf{Simp}$ of a simplicial complex K , we can construct a refinement L_\bullet by extending the partial order induced by the filtration on the simplices of K into a total order, and filter K according to this total order.

Using these two definitions, we can use the matrix reduction algorithm to compute the persistence barcode of any essentially finite filtration.

Definition 13. A simplicial filtration $G : \mathbb{R} \rightarrow \mathbf{Simp}$ is *finite* if it is essentially finite and the simplicial complex at any index is finite.

A persistence module $M : \mathbb{R} \rightarrow \mathbf{Vect}$ has *finite type* if it is essentially finite and the vector space at any index is finite-dimensional.

Let $(F_t)_{t \in \mathbb{R}}$ be a \mathbb{R} -indexed filtration of a finite simplicial complex K . Then there exists a simplexwise filtration $(K_i)_{1 \leq i \leq n}$ and a monotonic reindexing map $r : \mathbb{R} \rightarrow [n] = \{1, \dots, n\}$ such that $F_t = K_{r(t)}$: extend the partial order on K induced by F_t and the partial order \subseteq on K (face order) to a single total order on K , which defines the simplexwise filtration $(K_i)_i$. Now each complex F_t appears as some K_i in the simplexwise filtration, and this defines the map $r : t \mapsto i$.

Corollary 14. *The persistence barcode of the simplexwise filtration K_\bullet determines the persistence barcode of the filtration F_\bullet :*

$$H_*(F_\bullet) = \bigoplus_{\substack{I \in B(H_*(K_\bullet)) \\ r^{-1}(I) \neq \emptyset}} \mathbb{F}_\bullet^{r^{-1}(I)}$$

Note that the barcode of F_\bullet is a *multiset* of intervals in \mathbb{R} (either of the form $[a, b)$ or of the form $[a, \infty)$). Each interval corresponds to homology in a certain dimension.

Note that the collection $\{U_I : I \in B(H_*(L_\bullet))\}$ may contain empty intervals and multiple copies of the same interval. We may define the barcode of a general finite-type persistence module as a multiset of intervals. These multisets can be formalized either as a function $\mathcal{I}_{\mathbb{R}} \rightarrow \mathbb{N}$ determining the multiplicity of each interval, or as a set of pairs $(I, k) \in \mathcal{I}_{\mathbb{R}} \times \mathbb{N}$, so that multiple instances of the same interval can be distinguished by different indices k .