
Multidimensional Persistence Module Classification via Lattice-Theoretic Convolutions

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Abstract

Multiparameter persistent homology has been largely neglected as an input to machine learning algorithms. We consider the use of lattice-based convolutional neural network layers as a tool for the analysis of features arising from multiparameter persistence modules. We find that these show promise as an alternative to convolutions for the classification of multidimensional persistence modules.

1 Introduction

Persistent homology has the ability to discern both the global topology [10] and local geometry [5] of finite metric spaces (e.g. embedded weighted graphs, point clouds in \mathbb{R}^d) making it a befitting feature for the purposes of training a neural network. Single-dimensional homological persistence has drawn recent attention in deep learning [12, 21, 3]. This is, in part, due to a wide range of efficient software libraries [19, 11, 2] for computing persistent homology, as well as a growing cookbook of recipes for featurizing barcodes from single dimensional persistence, including persistence images [1], persistence landscapes [4], and more exotic methods [13].

Multidimensional persistence generalizes single-dimensional persistent homology in order to tackle filtrations parameterized in multiple dimensions. Unfortunately, there is no complete compact barcode-like characterization of multidimensional persistence modules [7]. We must make do with incomplete invariants. There are various algebraic invariants; in this paper we will use the *Hilbert function* and the *multi-graded Betti numbers*¹, both of which are \mathbb{N} -valued functions on the parameter space. The Hilbert function is nothing more than the pointwise (topological) Betti numbers, and the multi-graded Betti numbers have a geometric interpretation in terms of births and deaths [14]. The *rank invariant*, another invariant, is shown to be complete in the case of single dimensional persistence [7]; due to its difficulty to compute it is not considered here.

Multidimensional persistence suffers two hindrances to its usefulness in machine learning. First, software for computing multidimensional persistence is scarce. To the authors' knowledge, RIVET is the only available software for computing multidimensional persistence [16]; RIVET specializes to 2-dimensional persistence and is focused on interactive visualization rather than machine-interpretable output. Second, there has been only very recent and preliminary activity [23, 8] focusing on extraction of suitable features for multidimensional persistent homology. A full-fledged deep learning pipeline for multidimensional persistence waits in the wings.

We hope to ignite further interest in filling both of these gaps. In this paper, we propose a naive featurization of multidimensional persistence modules based on the aforementioned invariants, and design an architecture for classifying these persistence modules. This architecture employs a lattice-theoretic notion of convolution, thereby respecting the order relation of the parameters of

¹*Caveat lector*: the multi-graded (algebraic) Betti numbers need not be confused with the topological Betti numbers, i.e. the rank of homology.

the persistence module. We implement our model and compare the performance of our proposed lattice-convolutional architecture with a (simplified) standard convolutional architecture.

Related work

Very recent advances have been made to featurize multidimensional persistence modules via landscapes [23] and images [8]. Our naive approach at featurizing 2-dimensional persistence modules has the advantage of being readily computable with existing software [16]. Persistence modules supported on lattices are of increased interest. By leveraging Möbius inversion, these persistence modules are shown to generate a stable persistence diagram [17] as well as factor through a nice category with pseudo-inverses [15].

2 Background

Space constraints require that this section be laconic. For a primer on persistent homology, see [10, 6]; for multiparameter persistent homology, see [16]. An introduction to lattices may be found in [9].

2.1 Rips complexes and persistent homology

Let (\mathcal{M}, d) be a finite metric space. The *Vietoris-Rips complex* of \mathcal{M} at scale r is the abstract simplicial complex $\text{Rips}_r(\mathcal{M})$ whose simplices are subsets of \mathcal{M} of diameter at most r . There is a natural inclusion $\text{Rips}_r(\mathcal{M}) \rightarrow \text{Rips}_{r'}(\mathcal{M})$ for $r \leq r'$.

Applying the simplicial homology functor (with coefficients in a field k) H_i to $\text{Rips}_r(\mathcal{M})$ produces a sequence of vector spaces $PH_i(r)$. The inclusions $\text{Rips}_r(\mathcal{M}) \rightarrow \text{Rips}_{r'}(\mathcal{M})$ induce maps $PH_i(r) \rightarrow PH_i(r')$, producing the data of a *persistence module*. This structure can be compactly described as a functor from \mathbb{R} , viewed as a category via its standard order structure, to the category \mathbf{Vect}_k of vector spaces over k . The simplicity of the category \mathbb{R} gives these persistence modules simple structure: they decompose as direct sums of interval modules $I_{[a,b)}$, which have $I_{[a,b)}(r) = k$ for $a \leq r < b$ and zero otherwise. The maps are the identity where possible and the zero map otherwise.

2.2 Multiparameter persistence

The Rips construction produces a filtration of simplicial complexes from a finite metric space; it is natural to consider the behavior of the homology functor over a pair of coherent filtrations. Consider a finite metric space (\mathcal{M}, d) and a filtration function $\rho : \mathcal{M} \rightarrow \mathbb{R}$. This data specifies a bifiltration of simplicial complexes given by

$$\mathbb{X}_{r,t} = \text{Rips}_r\{x \in \mathcal{M} \mid \rho(x) \leq t\}.$$

There is a natural inclusion $\mathbb{X}_{r,t} \hookrightarrow \mathbb{X}_{r',t'}$ whenever $(r,t) \leq (r',t')$ in the lattice $\mathbb{R} \times \mathbb{R}$. Composing with the homology functor produces a 2-parameter persistence module

$$PH_i : \mathbb{R}_+ \times \mathbb{R} \rightarrow \mathbf{Vect}; \quad (r,t) \mapsto H_i(X_{r,t}).$$

While there does not exist a complete discrete invariant for PH_i , we can extract meaningful features. Two particularly informative types of features are the *Hilbert function*

$$\text{Hilb} : \mathbb{R}_+ \times \mathbb{R} \rightarrow \mathbb{Z}_+; \quad (r,t) \mapsto \dim(PH_i(r,t)),$$

and the *multi-graded Betti numbers* $\xi_j : \mathbb{R}_+ \times \mathbb{R} \rightarrow \mathbb{Z}_+$, for $j = 0, 1, 2$. For PH_i as above, the Hilbert function counts the number of connected components ($i = 0$), cycles ($i = 1$), or higher dimensional voids ($i > 1$) of the complex $\mathbb{X}_{r,t}$ at each $(r,t) \in \mathbb{R}_+ \times \mathbb{R}$. The multi-graded Betti numbers, on the other hand, capture information about births and deaths of homology classes.²

² $\xi_j(r,t)$ counts the rank of the j th term of the free resolution of PH_i at grading (r,t) [16].

2.3 Lattice-theoretic signal processing

Classical signal processing proceeds by constructing filters for space- or time-indexed signals using convolutional operators. These implicitly rely on the algebraic properties of the underlying space, in particular the existence of well-behaved translation operators. This fact is exploited in the general framework of algebraic signal processing [22] and extended to more general domains by the theory of graph signal processing [18]. We here describe a similar extension to signals on a finite lattice proposed in [20].

A *lattice* L is a partially ordered set in which every pair of elements x, y has a greatest lower bound (the *meet* $x \wedge y$) and a least upper bound (the *join* $x \vee y$). These operations and their properties produce an algebraic characterization of lattices; the ordering can be recovered from the algebra and vice versa. The key insight of [20] is that the meet and join operations on a lattice define two “shift operators” that can be exploited to define convolutional filters for signals on a lattice.

That is, for two signals $f, g : L \rightarrow \mathbb{R}$, where L is a lattice, we define

$$(f *_{\wedge} g)(x) = \sum_{a \in L} f(x \wedge a)g(a) \quad \text{and} \quad (f *_{\vee} g)(x) = \sum_{a \in L} f(x \vee a)g(a).$$

A nice class of examples of lattices are given by the sets \mathbb{R}^n viewed as partially ordered sets, with the ordering $(x_1, \dots, x_n) \leq (y_1, \dots, y_n)$ whenever $x_i \leq y_i$ for all $1 \leq i \leq n$. These are, of course, the indexing sets for persistent homology, suggesting that lattice convolutions over \mathbb{R}^n or its finite sublattices may be useful in processing data coming from multiparameter persistence computations.

3 Lattice Convolutional Neural Networks

Convolutions over \mathbb{R}^2 (with its abelian group structure) have served as an easily parameterized and efficient set of linear operations adapted to the structure of images. Their extreme utility in computer vision problems is owed to the translation equivariance properties of images: humans naturally recognize an image translated via an additive reparameterization as equivalent to the original.

The data of a multidimensional persistence module is also indexed by \mathbb{R}^n or a regular finite subset thereof, but its natural algebraic structure is not that of an abelian group. Rather, with its partial order structure, the indexing set is a lattice. In processing signals associated with a persistence module, it may be useful to take this structure into account rather than imposing the abelian group structure implied by standard convolutions.

To this end, we construct a lattice convolution-based neural network layer suitable for use with features originating from multidimensional persistence modules. To the authors’ knowledge, such an architecture has not previously been described, although a special case (where the underlying lattice is a power set) has been implemented in [24]. We specialize the convolutions described in Section 2.3 to the particular case of regular finite sublattices of \mathbb{R}^2 . These may be represented (up to isomorphism) as $L = [m] \times [n]$, where $[n]$ is the ordered set $\{0, 1, \dots, n\}$. The meet and join operations are easily computed elementwise:

$$(r, t) \wedge (r', t') = (\min(r, r'), \min(t, t')); \quad (r, t) \vee (r', t') = (\max(r, r'), \max(t, t')).$$

A lattice convolution layer takes as input an N_{in} -dimensional signal $f : [m] \times [n] \rightarrow \mathbb{R}^{N_{\text{in}}}$ and outputs an N_{out} -dimensional signal $[m] \times [n] \rightarrow \mathbb{R}^{N_{\text{out}}}$. The layer’s parameters are given by a function $g : [m] \times [n] \rightarrow \mathbb{R}^{N_{\text{out}} \times N_{\text{in}}}$. If we label the entries of $f(x, y)$ by f_i and the entries of $g(x, y)$ by g_j^i , the layer then acts by

$$\text{MeetConv}(f)(x, y)^j = \sum_i (f_i *_{\wedge} g_j^i)(x, y) = \sum_i \sum_{(a, b) \in [m] \times [n]} f_i(x \wedge a, y \wedge b) g_j^i(a, b)$$

in the case of convolution with respect to the meet operation, and

$$\text{JoinConv}(f)(x, y)^j = \sum_i (f_i *_{\vee} g_j^i)(x, y) = \sum_i \sum_{(a, b) \in [m] \times [n]} f_i(x \vee a, y \vee b) g_j^i(a, b)$$

in the case of convolution with respect to the join operation.

Traditional convolutional neural networks are useful in part because the convolution kernels (here the functions g) can have very small support, reducing the number of parameters that must be learned. In the standard convolutional setting, these kernels are implicitly supported in a neighborhood of the origin, but the location of the kernel is not usually explicitly specified. In the lattice setting, we do need to specify where the kernel resides. In the abelian group case, the kernel is supported near the identity, and similarly, when we treat our domain as a lattice, the kernel should be supported near the neutral element of the operation. That is, for a meet convolution, g should be supported at the maximum (m, n) , and for a join convolution, g should be supported at the minimum $(0, 0)$. This ensures that the convolution operators are capable of preserving information at every point of the space. For instance, if $g(0, 0) \neq 0$, then $(f *_\vee g)(x, y)$ is a sum of terms including $f(x, y)g(0, 0)$, so all information from one layer can be passed to the next layer.

4 Experiments

We use a small portion of the Princeton ModelNet dataset [25] as a source of finite metric spaces. This dataset consists of hundreds of 3-dimensional CAD models representing objects from 10 classes. We select two of the classes and sample points from the 3D models to produce finite metric spaces embedded in \mathbb{R}^3 . We then compute the corresponding multidimensional persistence modules, from which we produce features used as an input to a convolutional neural net classifier.

The pipeline thus begins with a 3D polyhedral model, of which 3000 vertices are sampled to produce a point cloud in \mathbb{R}^3 . This point cloud then produces a bifiltered simplicial complex, whose degree-0 persistent homology we calculate using RIVET [16], sampled at a discrete grid of 40×40 points, producing lattice-indexed signals given by the Hilbert function and the multi-graded Betti numbers ξ_0, ξ_1, ξ_2 ; four features in total. These are then passed to the classifier, which produces a class prediction, in this case either “sofa” or “monitor.”

As the filter function on these data sets, we use the *codensity* function

$$\rho_{\text{codense}}(x; k) = \left(\frac{1}{k} \sum_{y \in N_k(x)} d(x, y) \right)^{-1},$$

where $N_k(x)$ is the set of the k nearest neighbors to x ; we select $k = 100$. The name codensity is appropriate because the points in the densest regions of \mathcal{M} appear earlier in the filtration. A folk theorem is that the two-parameter persistent homology of a Rips/codensity bifiltration is stable under non-Hausdorff perturbations: *the (Rips) persistent homology of a point sample and another obtained by adding a small number of points at random are close with respect to the interleaving distance.*

We compare the performance of two convolutional networks on this classification task. One uses the lattice-convolution based layers described in Section 3, and the other uses standard convolutional layers. Each has three convolutional layers followed by three fully connected layers. The lattice-based convolution layers are of the form $\alpha \text{MeetConv}(x) + (1 - \alpha) \text{JoinConv}(x)$ for a hyperparameter $\alpha \in [0, 1]$. All convolution kernels have dimension 4×4 , hidden convolution layers have 16 features, and the final convolution has 8 features. The fully connected layers have 32 features at each inner layer. We use a cross entropy loss function with a softmax in the final layer. The networks are trained using the Adam gradient algorithm with learning rate 0.001 for 50 epochs. We reserve 10% of the data for testing. Results are shown in Figure 1.

Figure 1: Comparison training curves for the lattice neural network and standard convolutional neural network.

5 Discussion

Our proposed featurization for persistence modules is rather naive, but well adapted to the use of lattice convolutions as a data processing method. The lattice convolutional neural network shows promise as a method for classifying features arising from a multiparameter persistence module. The algebraic

perspective on partially ordered sets, exemplified by lattices, may also offer approaches to featurizing more complex invariants of persistence modules. In particular, the incidence algebra may offer a natural way to represent the rank invariant [7] in a way amenable to convolution-like operations. We hope with these brief experiments to inspire further work on featurizing multidimensional persistence for use in machine learning algorithms.

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