Persistent spectral graph

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Abstract

Persistent homology is constrained to purely topological persistence, while multiscale graphs account only for geometric information. This work introduces persistent spectral theory to create a unified low-dimensional multiscale paradigm for revealing topological persistence and extracting geometric shapes from high-dimensional datasets. For a point-cloud dataset, a filtration procedure is used to generate a sequence of chain complexes and associated families of simplicial complexes and chains, from which we construct persistent combinatorial Laplacian matrices. We show that a full set of topological persistence can be completely recovered from the harmonic persistent spectra, that is, the spectra that have zero eigenvalues, of the persistent combinatorial Laplacian matrices. However, non-harmonic spectra of the Laplacian matrices induced by the filtration offer another powerful tool for data analysis, modeling, and prediction. In this work, fullerene stability is predicted by using both harmonic spectra and non-harmonic persistent spectra, while the latter spectra are successfully devised to analyze the structure of fullerenes and model protein flexibility, which cannot be straightforwardly extracted from the current persistent homology. The proposed method is found to provide excellent predictions of the protein B-factors for which current popular biophysical models break down.

Keywords
persistent spectral analysis, persistent spectral graph, persistent spectral theory, spectral data analysis

1 | INTRODUCTION

Graph theory, a branch of discrete mathematics, concerns the relationship between objects. These objects can be either simple vertices, that is, nodes and/or points (zero simplexes), or high-dimensional simplexes. Here, the relationship refers to connectivity with possible orientations. Graph theory has many branches, such as geometric graph theory, algebraic graph theory, and topological graph theory. The study of graph theory draws on many other areas of mathematics, including algebraic topology, knot theory, algebra, geometry, group theory, combinatorics, and so on. For example, algebraic graph theory can be investigated by using either linear algebra, group theory, or graph invariants. Among them, the use of learning algebra in graph study leads to spectral graph theory (SGT).

Precursors of the spectral theory have often had a geometric flavor. An interesting spectral geometry question asked by Mark Kac was “Can one hear the shape of a drum?”¹ The Laplace-Beltrami operator on a closed Riemannian manifold has been intensively studied.² Additionally, eigenvalues and isoperimetric properties of graphs are the foundation of the explicit constructions of expander graphs.³ Moreover, the study of random walks
and rapidly mixing Markov chains utilized the discrete analog of the Cheeger inequality.\(^4\) The interaction between spectral theory and differential geometry became one of the critical developments.\(^2\) For example, the spectral theory of the Laplacian on a compact Riemannian manifold is a central object of de Rham-Hodge theory.\(^2\) Note that the Hodge Laplacian spectrum contains the topological information of the underlying manifold. Specifically, the harmonic part of the Hodge Laplacian spectrum corresponds to topological cycles. Connections between topology and SGT also play a central role in understanding the connectivity properties of graphs.\(^6-9\) Similarly, as the topological invariants revealing the connectivity of a topological space, the multiplicity of 0 eigenvalues of a 0-combinatorial Laplacian matrix is the number of connected components of a graph. Indeed, the number of \(q\)-dimensional holes can also be unveiled from the number of 0 eigenvalues of the \(q\)-combinatorial Laplacian.\(^10-13\) Nonetheless, SGT offers additional non-harmonic spectral information beyond topological invariants.

The traditional topology and homology are independent of metrics and coordinates and thus retain little geometric information. This obstacle hinders their practical applicability in data analysis. Recently, persistent homology has been introduced to overcome this difficulty by creating low-dimensional multiscale representations of a given object of interest.\(^14-19\) Specifically, a filtration parameter is devised to induce a family of geometric shapes for a given initial data. Consequently, the study of the underlying topologies or homology groups of these geometric shapes leads to the so-called topological persistence. Like the de Rham-Hodge theory that bridges differential geometry and algebraic topology, persistent homology bridges multiscale analysis and algebraic topology. Topological persistence is the most important aspect of the popular topological data analysis (TDA)\(^20-23\) and has had tremendous success in computational biology\(^24,25\) and worldwide competitions in computer-aided drug design.\(^26\)

Graph theory has been applied in various fields.\(^27\) For example, SGT is applied to the quantum calculation of \(\pi\)-delocalized systems. The Hückel method, or Hückel molecular orbital theory, describes the quantum molecular orbitals of \(\pi\)-electrons in \(\pi\)-delocalized systems in terms of a kind of adjacency matrix that contains atomic connectivity information.\(^28,29\) Additionally, the Gaussian network model (GNM)\(^30\) and anisotropic network model (ANM)\(^31\) represent protein \(C_\alpha\) atoms as an elastic mass-and-spring network by graph Laplacians. These approaches were influenced by the Flory theory of elasticity and the Rouse model.\(^32\) Like traditional topology, traditional graph theory extracts very limited information from data. In our earlier work, we have proposed multiscale graphs, called multiscale flexibility index (mFRI), to describe the multiscale nature of biomolecular interactions,\(^33\) such as hydrogen bonds, electrostatic effects, van der Waals interactions, hydrophilicity, and hydrophobicity. A multiscale spectral graph method has also been proposed as generalized GNM and generalized ANM.\(^34\) Our essential idea is to create a family of graphs with different characteristic length scales for a given dataset. We have demonstrated that our multiscale weighted colored graph (MWCG) significantly outperforms traditional spectral graph methods in protein flexibility analysis.\(^35\) More recently, we demonstrate that our MWCG outperforms other existing approaches in protein-ligand binding scoring, ranking, docking, and screening.\(^36\)

The objective of the present work is to introduce the persistent spectral graph as a new paradigm for the multiscale analysis of the topological invariants and geometric shapes of high-dimensional datasets. Motivated by the success of persistent homology\(^25\) and multiscale graphs\(^36\) in dealing with complex biomolecular data, we construct a family of spectral graphs induced by a filtration parameter. In the present work, we consider the radius filtration via the Vietoris-Rips complex, while other filtration methods can be implemented as well. As the filtration radius is increased, a family of persistent \(q\)-combinatorial Laplacians is constructed for a given point-cloud dataset. The diagonalization of these persistent \(q\)-combinatorial Laplacian matrices gives rise to persistent spectra. It is noted that our harmonic persistent spectra of 0-eigenvalues fully recover the persistent barcode or persistent diagram of persistent homology. Additional information is generated from non-harmonic persistent spectra, namely, the non-zero eigenvalues and associated eigenvectors. In combination with a simple machine learning algorithm, this additional spectral information is found to provide a powerful new tool for the quantitative analysis of molecular data, including the prediction of a set of protein B-factors for which existing standard predictors fail to work.

### 2 | THEORIES AND METHODS

In this section, we give a brief review of SGT and simplicial complex to establish notations and provide essential background. Subsequently, we introduce persistent spectral analysis.
2.1 Spectral graph theory

Graph structure encodes inter-dependencies among constituents and provides low-dimensional representations of high-dimensional datasets. One of the representations frequently used in SGT is to associate graphs with matrices, such as the Laplacian matrix and adjacency matrix. Analyzing the spectra from such matrices leads to the understanding of the topological and spectral properties of the graph.

Let $V$ be the vertex set, and $E$ be the edge set. For a given simple graph $G(V, E)$ (a simple graph can be either connected or disconnected), the degree of the vertex $v \in V$ is the number of edges that are adjacent to $v$, denoted $\text{deg}(v)$. The adjacency matrix $A$ is defined by

$$A(G) = \begin{cases} 1 & \text{if } v_i \text{ and } v_j \text{ are adjacent}, \\ 0 & \text{otherwise}. \end{cases}$$

and the Laplacian matrix $L$ is given by

$$L(G) = \begin{cases} \text{deg}(v_i) & \text{if } v_i = v_j, \\ -1 & \text{if } v_i \text{ and } v_j \text{ are adjacent}, \\ 0 & \text{otherwise}. \end{cases}$$

Obviously, the adjacency matrix characterizes the graph connectivity. The above two matrices are related through diagonal matrix $D$

$$L = D - A$$

Assuming $G(V, E)$ has $N$ nodes, then adjacency matrix $A$ and Laplacian matrix $L$ are both real symmetric $N \times N$ matrices. The eigenvalues of adjacency and Laplacian matrices are denoted and ordered as

$$\alpha_{\min} = \alpha_1 \leq \cdots \leq \alpha_2 \leq \alpha_1 = \alpha_{\max}$$

$$\lambda_{\min} = \lambda_1 \leq \lambda_2 \leq \cdots \leq \lambda_N = \lambda_{\max}.$$ (3)

The spectra of $A$ and $L$ have several interesting proprieties. To understand the robustness and connectivity of a graph, the algebraic connectivity $\lambda_2$ and the multiplicity of 0 eigenvalues are taken into consideration, which will be illustrated in Theorem 2 and Remark 1. For more detailed theorems and proofs, we refer the interested reader to a survey on Laplacian eigenvalues of graphs. Furthermore, some interesting properties and examples can be found in the Supporting Material S1 as well.

**Theorem 2.1.** Let $G(V, E)$ be a simple graph of order $N$, then the multiplicity of 0 eigenvalue for Laplacian matrix is the number of connected components of $G(V, E)$. The vertex degree is the value of the diagonal entry.

**Remark 2.1.** In this section, the Laplacian spectrum of simple graph $G(V, E)$ is concerned. For

$$G(V, E) = G_1(V_1, E_1) \cup \cdots \cup G_m(V_m, E_m), m \geq 1, m \in \mathbb{Z},$$

where $G_i(V_i, E_i) \subset G(V, E), i = 1, \cdots, m$ is a connected simple graph. If $m \geq 2$, the zero eigenvalue of $L(G)$ has multiplicity $m$, which results in algebraic connectivity $\lambda_2 = 0$. However, $\lambda_2 = 0$ cannot give any information about the $G_i(V_i, E_i)$. Therefore, we study the smallest non-zero eigenvalue of $L(G)$, which is actually the smallest algebraic connectivity of $L(G_i), i = 1, \cdots, m$. In Section 2.3 and Section 3, we analyze the smallest non-zero eigenvalue of the Laplacian matrix. To make the expression more concise, we still use $\hat{\lambda}_2$ as the smallest non-zero eigenvalue. If $G(V, E)$ is a connected simple graph, that is, $m = 1$, one has $\lambda_2 = \hat{\lambda}_2$. 

2.2 | Simplicial complex

A simplicial complex is a powerful algebraic topology tool that has a wide range of applications in graph theory, topological data analysis,17 and many physical fields.25 We briefly review simplicial complexes to generate notation and provide essential preparation for introducing persistent spectral graph.

2.2.1 | Simplex

Let \( \{v_0, v_1, \cdots, v_q\} \) be a set of points in \( \mathbb{R}^n \). A point \( v = \sum_{i=0}^{q} \lambda_i v_i \) is an affine combination of \( v_i \) if \( \sum_{i=0}^{q} \lambda_i = 1 \). An affine hull is the set of affine combinations. Here, \( q + 1 \) points \( v_0, v_1, \cdots, v_q \) are affinely independent if \( v_1 - v_0, v_2 - v_0, \cdots, v_q - v_0 \) are linearly independent. A \( q \)-plane is well defined if the \( q + 1 \) points are affinely independent. In \( \mathbb{R}^n \), one can have at most \( n \) linearly independent vectors. Therefore, there are at most \( n + 1 \) affinely independent points. An affine combination \( v = \sum_{i=0}^{q} \lambda_i v_i \) is a convex combination if all \( \lambda_i \) are non-negative. The convex hull is the set of convex combinations.

A \((\text{geometric})\) \( q \)-simplex denoted as \( \sigma_q \) is the convex hull of \( q + 1 \) affinely independent points in \( \mathbb{R}^n \) \((n \geq q)\) with dimension \( \dim(\sigma_q) = q \). A 0-simplex is a vertex, a 1-simplex is an edge, a 2-simplex is a triangle, and a 3-simplex is a tetrahedron, as shown in Figure 1. The convex hull of each nonempty subset of \( q + 1 \) points forms a subsimplex and is regarded as a face of \( \sigma_q \) denoted \( \tau \).

2.2.2 | Simplicial complex

A (finite) simplicial complex \( K \) is a (finite) collection of simplices in \( \mathbb{R}^n \) satisfying the following conditions.

1. If \( \sigma_q \in K \) and \( \sigma_p \) is a face of \( \sigma_q \), then \( \sigma_p \in K \).
2. The non-empty intersection of any two simplices \( \sigma_q, \sigma_p \in K \) is a face of both \( \sigma_q \) and \( \sigma_p \).

Each element \( \sigma_q \in K \) is a \( q \)-simplex. The dimension of \( K \) is defined as \( \dim(K) = \max\{\dim(\sigma_q) : \sigma_q \in K \} \). To distinguish topological spaces based on the connectivity of simplicial complexes, one uses Betti numbers. The \( k \)th Betti number, \( \beta_k \), counts the number of \( k \)-dimensional holes on a topological surface. The geometric meaning of Betti numbers in \( \mathbb{R}^3 \) is the following: \( \beta_0 \) represents the number of connected components, \( \beta_1 \) counts the number of one-dimensional loops or circles, and \( \beta_2 \) describes the number of two-dimensional voids or holes. In a nutshell, the Betti number sequence \( \{\beta_0, \beta_1, \beta_2, \cdots\} \) reveals the intrinsic topological property of the system.

Recall that in graph theory, the degree of a vertex (0-simplex) \( v \) is the number of edges that are adjacent to the vertex, denoted as \( \deg(v) \). However, once we generalize this notion to \( q \)-simplex, problem aroused since \( q \)-simplex can have \((q - 1)\)-simplices and \((q + 1)\)-simplices adjacent to it at the same time. Therefore, the upper adjacency and lower adjacency are required to define the degree of a \( q \)-simplex for \( q > 0 \).

**Definition 2.1.** Two \( q \)-simplices \( \sigma_q^i \) and \( \sigma_q^j \) of a simplicial complex \( K \) are lower adjacent if they share a common \((q - 1)\)-face, denoted \( \sigma_q^i \sim \sigma_q^j \). The lower degree of \( q \)-simplex, denoted \( \deg_L(\sigma_q) \), is the number of nonempty \((q - 1)\)-simplices in \( K \) that are faces of \( \sigma_q \) which is always \( q + 1 \).

![Figure 1](image-url)  
**Figure 1** Illustration of simplices. A, 0-simplex (a vertex); B, 1-simplex (an edge); C, 2-simplex (a triangle); and D, 3-simplex (a tetrahedron)
Definition 2.2. Two $q$-simplices $\sigma_q^i$ and $\sigma_q^j$ of a simplicial complex $K$ are upper adjacent if they share a common $(q+1)$-face, denoted $\sigma_q^i \sim \sigma_q^j$. The upper degree of $q$-simplex, denoted $\deg_U(\sigma_q)$, is the number of $(q+1)$-simplices in $K$ of which $\sigma_q$ is a face.

Then, the degree of a $q$-simplex ($q > 0$) is defined as:

$$\deg(\sigma_q) = \deg_L(\sigma_q) + \deg_U(\sigma_q) = \deg_U(\sigma_q) + q + 1.$$  \hspace{1cm} (4)

A supplemental example that illustrates the relation between simplicial complex and its corresponding Betti number can be found in the Supporting Material S2.

2.2.3 | Chain complex

Chain complex is an important concept in topology, geometry, and algebra. Let $K$ be a simplicial complex of dimension $q$. A $q$-chain is a formal sum of $q$-simplices in $K$ with $\mathbb{Z}_2$ field of the coefficients for the sum. Under the addition operation of $\mathbb{Z}_2$, a set of all $q$-chains is called a chain group and denoted as $C_q(K)$. To relate these chain groups, we denote boundary operator by $\partial_q : C_q(K) \rightarrow C_{q-1}(K)$. The boundary operator maps a $q$-chain which is a linear combination of $q$-simplices to the same linear combination of the boundaries of the $q$-simplices. Denoting $\sigma_q = [v_0, v_1, \cdots, v_q]$ for the $q$-simplex spanned by its vertices, its boundary operator can be defined as:

$$\partial_q \sigma_q = \sum_{i=0}^{q} (-1)^i \sigma_{q-1}^i,$$  \hspace{1cm} (5)

with $\sigma_q = [v_0, \cdots, v_q]$ being the $q$-simplex. Here, $\sigma_{q-1}^i = [v_0, \cdots, \hat{v}_i, \cdots, v_q]$ is the $(q-1)$-simplex with $v_i$ being omitted. A $q$-chain is called $q$-cycle if its boundary is zero. A chain complex is the sequence of chain groups connected by boundary operators

$$\cdots \xrightarrow{\partial_{q+1}} C_{q+1}(K) \xrightarrow{\partial_q} C_q(K) \xrightarrow{\partial_{q-1}} C_{q-1}(K) \rightarrow \cdots$$  \hspace{1cm} (6)

2.3 | Persistent spectral analysis

In this section, we introduce persistent spectral theory (PST) to extract rich topological and spectral information of simplicial complexes via a filtration process. We briefly review preliminary concepts about the oriented simplicial complex and $q$-combinatorial Laplacian, while more detail information can be found elsewhere.\textsuperscript{11-13,37} Then, we discuss the properties of the $q$-combinatorial Laplacian matrix together with its spectrum. Moreover, we employ the $q$-combinatorial Laplacian to establish the PST. Finally, we discuss some variants of the persistent $q$-combinatorial Laplacian matrix and illustrate their formulation on simple geometry, that is, a benzene molecule.

2.3.1 | Oriented simplicial complex and $q$-combinatorial Laplacian

An oriented simplicial complex is the one in which all of the simplices in the simplicial complex, except for vertices and $\emptyset$, are oriented. A $q$-combinatorial Laplacian is defined based on oriented simplicial complexes, and its lower- and higher-dimensional simplexes can be employed to study a specifically oriented simplicial complex.

We first introduce oriented simplex complexes. Let $\sigma_q$ be a $q$-simplex, we can define the ordering of its vertex set. If two orderings defined on $\sigma_q$ differ from each other by an even permutation, we say that they are equivalent, and each of them is called an orientation of $\sigma_q$. An oriented $q$-simplex is a simplex $\sigma_q$ with the orientation of $\sigma_q$. An oriented
simplicial complex $K$ is defined if all of its simplices are oriented. Suppose $\sigma^i_q$ and $\sigma^j_q \in K$ with $K$ being an oriented simplicial complex. If $\sigma^i_q$ and $\sigma^j_q$ are upper adjacent with a common upper $(q+1)$-simplex $\tau_{q+1}$, we say they are similarly oriented if both have the same sign in $\partial_q(\tau_{q+1})$ and dissimilarly oriented if the signs are opposite. Additionally, if $\sigma^i_q$ and $\sigma^j_q$ are lower adjacent with a common lower $(q-1)$-simplex $\eta_{q-1}$, we say they are similarly oriented if $\eta_{q-1}$ has the same sign in $\partial_q(\sigma^i_q)$ and $\partial_q(\sigma^j_q)$, and dissimilarly oriented if the signs are opposite.

Similarly, we can define $q$-chains based on an oriented simplicial complex $K$. The $q$-chain $C_q(K)$ is also defined as the linear combinations of the basis, with the basis being the set of oriented $q$-simplices of $K$. The $q$-boundary operator $\partial_q : C_q(K) \to C_{q-1}(K)$ is

$$\partial_q \sigma_q = \sum_{i=0}^{q} (-1)^i \sigma^i_{q-1}, \quad (7)$$

with $\sigma_q = [v_0, \ldots, v_q]$ to be the oriented $q$-simplex, and $\sigma^i_{q-1} = [v_0, \ldots, \hat{v}_i, \ldots, v_q]$ the oriented $(q-1)$-simplex with its vertex $v_i$ being removed. Let $\partial_q$ be the matrix representation of a $q$-boundary operator with respect to the standard basis for $C_q(K)$ and $C_{q-1}(K)$ with some assigned orderings. Then, the number of rows in $\partial_q$ corresponds to the number of $(q-1)$-simplices and the number of columns shows the number of $q$-simplices in $K$, respectively. Associated with the $q$-boundary operator is the adjoint operator denoted $q$-adjoint boundary operator, defined as

$$\partial^*_q : C_{q-1}(K) \to C_q(K), \quad (8)$$

and the transpose of $\partial_q$, denoted $B^T_q$, is the matrix representation of $\partial^*_q$ relative to the same ordered orthonormal basis as $\partial_q$.

Let $K$ be an oriented simplicial complex, for integer $q \geq 0$, the $q$-combinatorial Laplacian is a linear operator $\Delta_q : C_q(K) \to C_q(K)$

$$\Delta_q = \partial_q \partial_q^* + \partial^* \partial_q, \quad (9)$$

with $\partial_q \partial_q^* = 0$, which implies $\text{Im}(\partial_q + 1) \subseteq \ker(\partial_q)$. The $q$-combinatorial Laplacian matrix, denoted $\mathcal{L}_q$, is the matrix representation of $\Delta_q$.

$$\mathcal{L}_q = B^U_q + B^L_q B^T_q B_q, \quad (10)$$

of operator $\Delta_q$, with $\mathcal{B}_q$ and $\mathcal{B}_{q+1}$ being the matrices of dimension $q$ and $q+1$. Additionally, we denote upper and lower $q$-combinatorial Laplacian matrices by $\mathcal{L}^U_q = B^U_{q+1} B^T_{q+1}$ and $\mathcal{L}^L_q = B^T_q B_q$, respectively. Note that $\partial_0$ is the zero map, which leads to $\mathcal{B}_0$ being a zero matrix. Therefore, $\mathcal{L}_0(K) = B^U_0 + B^L_0 B_0$, with $K$ the (oriented) simplicial complex of dimension 1, which is actually a simple graph. In particular, 0-combinatorial Laplacian matrix $\mathcal{L}_0(K)$ is actually the Laplacian matrix defined in the SGT. In fact, Equation (2) is exactly the same as Equation (14) given in the following.

Given an oriented simplicial complex $K$ with $0 \leq q \leq \dim(K)$, one can obtain the entries of its corresponding upper and lower $q$-combinatorial Laplacian matrices explicitly.

$$\left( \mathcal{L}^U_q \right)_{ij} = \begin{cases} \deg_U(\sigma^i_q), & \text{if } i = j, \\ 1, & \text{if } i \neq j, \sigma^i_q \sim \sigma^j_q \text{ with similar orientation}, \\ -1, & \text{if } i \neq j, \sigma^i_q \sim \sigma^j_q \text{ with dissimilar orientation}, \\ 0, & \text{otherwise}. \end{cases} \quad (11)$$
The entries of $q$-combinatorial Laplacian matrices are

$$q > 0, \ (L_q)_{ij} = \begin{cases} \deg(\sigma_q^i) + q + 1, & \text{if } i = j. \\ 1, & \text{if } i \neq j, \sigma_q^i \not\sim \sigma_q^j \text{ with similar orientation.} \\ -1, & \text{if } i \neq j, \sigma_q^i \not\sim \sigma_q^j \text{ with dissimilar orientation.} \\ 0, & \text{otherwise.} \end{cases}$$

(13)

$$q = 0, \ (L_q)_{ij} = \begin{cases} \deg(\sigma_0^i), & \text{if } i = j. \\ -1, & \text{if } \sigma_0^i \not\sim \sigma_0^j. \\ 0, & \text{otherwise.} \end{cases}$$

(14)

### 2.3.2 Spectral analysis of $q$-combinatorial Laplacian matrices

A $q$-combinatorial Laplacian matrix for oriented simplicial complexes is a generalization of the Laplacian matrix in graph theory. The spectra of a Laplacian matrix play an essential role in understanding the connectivity and robustness of simple graphs (simplicial complexes of dimension 1). They can also distinguish different topological structures. Inspired by the capability of the Laplacian spectra of analyzing topological structures, we study the spectral properties of $q$-combinatorial Laplacian matrices to reveal topological and spectral information of simplicial complexes with dimension $0 \leq q \leq \dim(K)$.

We clarify that for a given finite simplicial complex $K$, the spectra of its $q$-combinatorial Laplacian matrix are independent of the choice of the orientation for the $q$-simplices of $K$. The proof can be found in Reference 13. Figure 2 provides a simple example to illustrate this property. In Figure 2, we have two oriented simplicial complexes, $K_1$ and $K_2$, with the same geometric structure but different orientations. For the sake of brevity, we use 1, 2, 3, 4, and 5 to represent 0-simplices (vertices); 12, 23, 34, 24, and 45 to describe 1-simplices (edges); and 234 to stand for the 2-simplex (triangle). Then the 0-combinatorial Laplacian matrix of $K_1$ and $K_2$ is

![Figure 2](image.png)

**Figure 2** Illustration of two oriented simplicial complexes with the same geometric structure but having different orientations. Here, we denote the vertices by 1, 2, 3, 4, and 5; edges by 12, 23, 34, 24, and 45; and the triangle by 234.
Obviously, $\mathcal{L}_0(K_1)$ and $\mathcal{L}_0(K_2)$ have the same spectra. For $q = 1$, there are five 1-simplices in $K_1$ and $K_2$, while 1-combinatorial Laplacian matrices have dimension $5 \times 5$. Using $K_1$ as an example, since 12 and 23 are lower adjacent with similar orientation, the element of $\mathcal{L}_1(K_1)$ addressed at first row and second column is 1 according to Equation (13). Since 34 and 45 are lower adjacent with dissimilar orientation, $(\mathcal{L}_1(K_1))_{35} = -1$. Moreover, 23 and 34 are upper adjacent, which results in $(\mathcal{L}_1(K_1))_{23} = 0$. For the diagonal parts, 12 and 45 are not the faces of any 2-simplex, while 23, 34, and 24 are the faces of 2-simplex 234. Therefore, $\deg U(12) = \deg U(12) = 0$ and $\deg U(23) = \deg U(34) = \deg U(34) = 1$, so the diagonal terms of $\mathcal{L}_0(K_1)$ are 2, 3, 3, 3, and 2.

The spectra of $\mathcal{L}_1(K_1)$ and $\mathcal{L}_1(K_2)$ have the same eigenvalues: $\left\{3, \frac{\sqrt{2} + \sqrt{5}}{2}, \frac{\sqrt{13} + \sqrt{2}}{2}\right\}$. Since $K_1$ and $K_2$ do not have 3-simplices, $\mathcal{L}_q$ is a zero matrix when $q \geq 3$.

A $q$-combinatorial Laplacian matrix is symmetric and positive semi-definite. Therefore, its eigenvalues are all real and non-negative. An analogy to the property that the number of zero eigenvalues of $\mathcal{L}_0$ represents the number of connected components ($\beta_0$) in the simple graph (simplicial complex with dimension 1), the number of zero eigenvalues of $\mathcal{L}_q$ can also reveal the topological information. More specifically, for a given finite-oriented simplicial complex, the Betti number $\beta_q$ of $K$ satisfies

$$\beta_q = \dim(\mathcal{L}_q(K)) - \text{rank}(\mathcal{L}_q(K)) = \text{nullity}(\mathcal{L}_q(K)) = \#\text{of zero eigenvalues of } \mathcal{L}_q(K). \quad (15)$$

In the Supporting material S3, we illustrate the connection between Betti number and the dimension of the rank of $q$-combinatorial Laplacian matrix.

### 2.3.3 Persistent spectral theory

Instead of using the aforementioned spectral analysis for $q$-combinatorial Laplacian matrix to describe a single configuration, we propose a PST to create a sequence of simplicial complexes induced by varying a filtration parameter, which is inspired by persistent homology and our earlier work in multiscale graphs.\(^{34,35}\) We provide a brief introduction to persistent homology in the Supporting Material S4.

A filtration of an oriented simplicial complex $K$ is a sequence of sub-complexes $(K_i)_{i=0}^m$ of $K$

$$\emptyset = K_0 \subseteq K_1 \subseteq K_2 \subseteq \cdots \subseteq K_m = K. \quad (16)$$

It induces a sequence of chain complexes
\[
\cdots C_{q+1} = \frac{\partial^{2}_{q+1}}{\partial q+1^2} C_q = \frac{\partial^{2}_{q}}{\partial q^2} \cdots = \frac{\partial^{2}_{1}}{\partial 1^2} C_2 = \frac{\partial^{2}_{2}}{\partial 2^2} C_1 = \frac{\partial^{2}_{3}}{\partial 3^2} C_0 = \frac{\partial^{2}_{0}}{\partial 0^2} C_{-1}
\]

where \( C_q = C_q(K_t) \) and \( \partial_q^t : C_q(K_t) \rightarrow C_{q-1}(K_t) \). Each \( K_t \) itself is an oriented simplicial complex, which has dimension denoted by \( \text{dim}(K_t) \). If \( q < 0 \), then \( C_q(K_t) = \{ \emptyset \} \) and \( \partial_q^t \) is actually a zero map. \(^2\) For a general case of \( 0 < q \leq \text{dim}(K_t) \), if \( \sigma_q \) is an oriented \( q \)-simplex of \( K_t \), then

\[
\partial_q^t(\sigma_q) = \sum_{i=0}^{q} (-1)^i \sigma_{q-i}, \sigma_q \in K_t,
\]

with \( \sigma_q = [v_0, \ldots, v_q] \) being the oriented \( q \)-simplex, and \( \sigma_{q-1} = [v_0, \ldots, \hat{v}_i, \ldots, v_q] \) being the oriented \((q-1)\)-simplex for which its vertex \( v_i \) is removed.

Let \( C_q^{t+p} \) be the subset of \( C_q^t \) whose boundary is in \( C_{q-1}^t \):

\[
C_q^{t+p} = \left\{ \alpha \in C_q^t \mid \partial^{t+p}_q(\alpha) \in C_{q-1}^t \right\}.
\]

We define

\[
\partial^{t+p}_q : C_q^{t+p} \rightarrow C_{q-1}^t.
\]

Based on the \( q \)-combinatorial Laplacian operator, the \( p \)-persistent \( q \)-combinatorial Laplacian operator \( \Delta^{t+p}_q : C_q(K_t) \rightarrow C_q(K_t) \) defined along the filtration can be expressed as

\[
\Delta^{t+p}_q = \partial^{t+p}_{q+1} \left( \partial^{t+p}_{q+1} \right)^T + \partial^{t+p}_q \partial^{t+p}_q.
\]

We denote the matrix representations of boundary operator \( \partial^{t+p}_{q+1} \) and \( \partial^{t+p}_q \) by \( B^{t+p}_{q+1} \) and \( B^{t+p}_q \), respectively. It is clear that the number of rows in \( B^{t+p}_{q+1} \) is the number of oriented \( q \)-simplices in \( K_t \), and the number of columns is the number of oriented \((q+1)\)-simplices in \( K_t \cap C_{q+1} \). The transpose values of the matrices \( B^{t+p}_{q+1} \) and \( B^{t+p}_q \) are the matrix representations of the adjoint boundary operator \( \left( \partial^{t+p}_{q+1} \right)^T \) and \( \partial^{t+p}_q \), respectively. Therefore, the \( p \)-persistent \( q \)-combinatorial Laplacian matrix, \( L^{t+p}_q \), is

\[
L^{t+p}_q = B^{t+p}_{q+1} \left( B^{t+p}_{q+1} \right)^T + \left( B^{t+p}_q \right)^T B^{t+p}_q.
\]

Intuitively, for a non-empty set \( C_q^t \), the \( p \)-persistent \( q \)-combinatorial Laplacian matrix \( L^{t+p}_q \) is a square matrix with dimension to be the number of \( q \)-simplices in \( K_t \). Moreover, \( L^{t+p}_q \) is symmetric and positively semi-defined and thus, all the spectra of \( L^{t+p}_q \) are real and non-negative. If \( p = 0 \), then \( L^{t+p}_q \) is exactly the \( q \)-combinatorial Laplacian matrix defined in Equation (10).

We are interested in the difference between \( L^{t+0}_q \) and \( L^{t+p}_q \). Suppose we have an oriented simplicial complex \( K_t \), and also an oriented simplicial complex \( K_{t+p} \) constructed by adding different dimension simplices ("outer" topological structures) to \( K_t \) with \( \text{dim}(K_{t+p}) = q+1 \). Since \( K_t \subset K_{t+p} \), we have
\[ L_{q}^{t+0} = B_{q+1}^{t+0} \left( B_{q+1}^{t+0} \right)^T + \left( B_{q}^{t} \right)^T B_{q}^{t+0} \]
\[ L_{q}^{t+p} = B_{q+1}^{t+p} \left( B_{q+1}^{t+p} \right)^T + \left( B_{q}^{t} \right)^T B_{q}^{t+p} \]

Case 1. If \( \text{Im}(\delta_{q+1}^{t+p}) \cap (K_{t+p} \setminus K_{t}) \cap C_{q} = \emptyset \) for all possible \( q \), then the “outer” topological structures are disconnected with \( K_{t} \). Therefore, the boundary matrix \( B_{q+1}^{t+p} \) is exactly the same as \( B_{q+1}^{t} \). In this situation, the spectra of \( L_{q}^{t+0} \) and \( L_{q}^{t+p} \) are the same, which reveals the fact that the topological structure \( K_{t} \) does not change under the filtration process.

Case 2. If \( \text{Im}(\delta_{q+1}^{t+p}) \cap (K_{t+p} \setminus K_{t}) \cap C_{q} \neq \emptyset \) for at least one \( q \), then the boundary matrix \( B_{q+1}^{t+p} \) will be changed by adding additional non-zero columns. Therefore, \( L_{q}^{t+p} \) is no longer the same as \( L_{q}^{t+0} \), but the structure information in \( K_{t} \) will still be preserved in \( L_{q}^{t+p} \). In this case, the topological structure \( K_{t} \) builds connection with “outer” topological structures. By calculating the spectra of \( L_{q}^{t+p} \), the disappeared and preserved structure information of \( K_{t} \) under the filtration process can be revealed.

Based on the fact that the topological and spectral information of \( K_{t} \) can also be analyzed from \( L_{q}(K_{t}) \) along with the filtration parameter by diagonalizing the \( q \)-combinatorial Laplacian matrix, we focus on the spectra information calculated from \( L_{q}^{t+p} \). Denote the set of spectra of \( L_{q}^{t+p} \) by

\[ \text{Spectra}(L_{q}^{t+p}) = \{(\lambda_1)_{q}^{t+p}, (\lambda_2)_{q}^{t+p}, \cdots, (\lambda_N)_{q}^{t+p}\}, \]

where \( L_{q}^{t+p} \) has dimension \( N \times N \) and spectra are arranged in ascending order. The smallest non-zero eigenvalue of \( L_{q}^{t+p} \) is defined as \( (\lambda_1)_{q}^{t+p} \). In the previous section, we have seen that Betti numbers (ie, \# of zero eigenvalues) can reveal \( q \)-cycle information. Similarly, we define the number of zero eigenvalues of \( p \)-persistent \( q \)-combinatorial Laplacian matrix \( L_{q}^{t+p} \) to be the \( p \)-persistent \( q \)th Betti numbers

\[ \beta_q^{t+p} = \dim(L_{q}^{t+p}) - \text{rank}(L_{q}^{t+p}) = \text{nullity}(L_{q}^{t+p}) = \# \text{ of zero eigenvalues of } L_{q}^{t+p}. \quad (22) \]

In fact, \( \beta_q^{t+p} \) counts the number of \( q \)-cycles in \( K_{t} \) that are still alive in \( K_{t+p} \), which exactly provides the same topological information as persistent homology does. However, PST offers additional geometric information from the spectra of persistent combinatorial Laplacian matrix beyond topological persistence. In general, the topological changes can be read off from persistent Betti numbers (harmonic persistent spectra) and the geometric changes can be derived from the non-harmonic persistent spectra.

Figure 3 demonstrates an example of a standard filtration process. Here the initial setup \( K_{1} \) consists of five 0-simplices (vertices). We construct Vietoris-Rips complexes by using an ever-growing circle centered at each vertex with radius \( r \). Once two circles overlapped with each other, a 1-simplex (edge) is formed. A 2-simplex (triangle) will be created when 3 circles contact with one another, and a 3-simplex will be generated once 4 circles get overlapped one another. As shown in Figure 3, we can attain a series of simplicial complexes from \( K_{1} \) to \( K_{6} \) with the radius of circles increasing. Table 1 lists the number of \( q \)-cycles of simplicial complex to fully illustrate how to construct \( p \)-persistent \( q \)-combinatorial Laplacian matrices by the boundary operator and determine persistent Betti numbers, we analyze 6 \( p \)-persistent \( q \)-combinatorial Laplacian matrices and their corresponding harmonic persistent spectra (ie, persistent Betti numbers) and non-harmonic persistent spectra. A supplementary example to distinguish different topological structures by implementing PST is provided in the Supporting Material S5. Moreover, additional matrices are analyzed in the Supporting Material S6.

Case 1 The initial setup is \( K_{3} \) and the end status is \( K_{4} \). The 1-persistent 0, 1, 2-combinatorial Laplacian operators are
Δ_3^1 = \partial_3^1 (\partial_3^1 + 1)^* + \partial_3^1 \partial_3^1,
Δ_1^2 = \partial_2^1 (\partial_2^1 + 1)^* + \partial_2^1 \partial_1^3,
Δ_2^3 = \partial_2^3 (\partial_2^3 + 1)^* + \partial_2^1 \partial_2^3,

Since 2-simplex and 3-simplex do not exist in K_4, \partial_2^3, \partial_1^3, and \partial_2^1 do not exist, then
\[ L_3^{1+1} = B_1^{1+1} (B_1^{1+1})^T + (B_0^3)^T B_0^3, \]
\[ L_1^{3+1} = (B_1^3)^T B_1^3. \]

From Table 2, one can see that \( \beta_0^{3+1} = 0 \) and \( \beta_1^{3+1} = 1 \), which reveals that only one 0-cycle and one 1-cycle in K_3 are still alive in K_4.

**Case 2** The initial setup is K_6 and the end status is K_6. The 0-persistent 0, 1, 2-combinatorial Laplacian operators are
\[ L_0^{6+0} = B_1^6 (B_1^6 + 0)^T + (B_0^6)^T B_0^6, \]
\[ L_1^{5+0} = B_2^5 (B_2^5 + 0)^T + (B_1^5)^T B_1^5, \]
\[ L_2^{3+0} = B_3^3 (B_3^3 + 0)^T + (B_2^3)^T B_2^3. \]

From Table 3, \( \beta_0^{6+0} = 1, \beta_1^{6+0} = 0 \), and \( \beta_2^{6+0} = 0 \) imply that only one 0-cycle (connected component) exists in K_6, with
We have constructed a family of persistent spectral graphs induced by a filtration parameter. For the sake of simplicity, we focus on the analysis of high-dimensional spectra with \( p = 0 \) in the rest of this section. As clarified before, the 0-persistent \( q \)-combinatorial Laplacian matrix is the \( q \)-combinatorial Laplacian matrix.

A graph structure encodes inter-dependencies among constituents and provides a convenient representation of the high-dimensional data. Naturally, the same idea can be applied to higher-dimensional spaces. For a set of points \( V \subset \mathbb{R}^n \) without additional structures, we consider growing an \((n-1)\)-sphere centered at each point with an ever-increasing radius \( r \). Therefore, a family of 0-persistent \( q \)-combinatorial Laplacian matrices as well as spectra can be generated as the radius \( r \) increases, which provides topological and spectral features to distinguish individual entries of the dataset.

### Variants of \( p \)-persistent \( q \)-combinatorial Laplacian matrices

The traditional approach in defining the \( q \)-boundary operator \( \partial_q : C_q(K) \rightarrow C_{q-1}(K) \) can be expressed as:

\[
\begin{align*}
\partial^1_q &= \begin{bmatrix}
0 & 0 & 1 & 0 & 0 & 0 & 0 & 0
\end{bmatrix}, & \partial^2_q &= \begin{bmatrix}
0 & 0 & 0 & 1 & 0 & 0 & 0 & 0
\end{bmatrix}.
\end{align*}
\]

\[
\begin{array}{cccc}
q & q = 0 & q = 1 & q = 2 \\
\hline
\partial^1_q & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} \\
\partial^2_q & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} \\
\partial^3_q & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} \\
\partial^4_q & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} \\
\partial^5_q & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} \\
\partial^6_q & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} & \begin{bmatrix}
0 & 1 & 2 & 3 & 4
\end{bmatrix} \\
\end{array}
\]
which leads to the corresponding elements in the boundary matrices being either 1 or −1. However, to encode more geometric information into the Laplacian operator, we add volume information of \(q\)-simplex \(\sigma_q\) to the expression of \(q\)-boundary operator.

Given a vertex set \(V = \{v_0, v_1, \cdots, v_q\}\) with \(q + 1\) isolated points (0-simplices) randomly arranged in the \(n\)-dimensional Euclidean space \(\mathbb{R}^n\), often with \(n \geq q\). Set \(d_{ij}\) to be the distances between \(v_i\) and \(v_j\) with \(0 \leq i \leq j \leq q\) and obviously, \(d_{ij} = d_{ji}\). The Cayley-Menger determinant can be expressed as\(^{39}\)

\[
\partial_q \sigma_q = \sum_{i=0}^{q} (-1)^i \sigma_{q-i},
\]
The $q$-dimensional volume of $q$-simplex $\sigma_q$ with vertices $\{v_0, v_1, \ldots, v_q\}$ is defined by

$$\text{Vol}(\sigma_q) = \sqrt{\frac{(-1)^{q+1}}{(q!)^2 2^q} \text{Det}_{CM}(v_0, v_1, \ldots, v_q)}.$$

In trivial cases, $\text{Vol}(\sigma_0) = 1$, meaning the 0-dimensional volume of 0-simplex is 1, that is, there is only 1 vertex in a 0-simplex. Also, the 1-dimensional volume of 1-simplex $\sigma_1 = [v_i, v_j]$ is the distance between $v_i$ and $v_j$, and the 2-dimensional volume of 2-simplex is the area of a triangle $[v_i, v_j, v_k]$.

In applications, it is often useful to replace the characteristic number “1” with some other descriptive quantities.\(^{40}\) Therefore, we define weighted boundary operator equipped with volume, denoted $\tilde{\partial}_q$,

$$\tilde{\partial}_q \sigma_q = \sum_{i=0}^{q} (-1)^i \text{Vol}(\sigma_q^i) \sigma_q^{i-1}. \quad (25)$$

Employing the same concept to the PST, we have the volume-weighted $p$-persistent $q$-combinatorial Laplacian operator. We also define

$$\tilde{\partial}_q^{t+p} : C_q^{t+p} \to C_q^{t-1} \quad (26)$$

with

$$C_q^{t+p} = \left\{ \alpha \in C_q^{t+p} \mid \tilde{\partial}_q^{t+p}(\alpha) \in C_q^{t-1} \right\}.$$

Similarly, an inverse-volume weighted boundary operator, denoted $\partial_q$, is given by

$$\partial_q \sigma_q = \sum_{i=0}^{q} (-1)^i \frac{1}{\text{Vol}(\sigma_q)} \sigma_q^{i-1}. \quad (27)$$

To define an inverse-volume weighted $p$-persistent $q$-combinatorial Laplacian operator. We define

$$\partial_q^{t+p} : C_q^{t+p} \to C_q^{t-1} \quad (28)$$

with

$$C_q^{t+p} = \left\{ \alpha \in C_q^{t+p} \mid \partial_q^{t+p}(\alpha) \in C_q^{t-1} \right\}.$$
Then volume-weighted and inverse-volume-weighted $p$-persistent $q$-combinatorial Laplacian operators defined along the filtration can be expressed as

$$
\Delta_{q}^{t+p} = \partial_{q+1}^{t+p} (\partial_{q+1}^{t+p})^* + \partial_{q}^{t} \partial_{q}^{t},
$$

$$
\Delta_{q}^{t+p} = \partial_{q+1}^{t+p} (\partial_{q}^{t+p})^* + \partial_{q}^{t} \partial_{q}^{t}.
$$

The corresponding weighted matrix representations of boundary operators $\partial_{q+1}^{t+p}, \partial_{q}^{t}, \partial_{q+1}^{t+p}$, and $\partial_{q}$ are denoted $B_{q+1}^{t+p}$, $B_{q}^{t}$, $B_{q+1}^{t+p}$, and $B_{q}$, respectively. Therefore, volume-weighted and inverse-volume-weighted $p$-persistent $q$-combinatorial Laplacian matrices can be expressed as

$$
\mathcal{L}_{q}^{t+p} = B_{q+1}^{t+p} (B_{q+1}^{t+p})^T + (B_{q}^{t})^T (B_{q}^{t}),
$$

$$
\mathcal{L}_{q}^{t+p} = B_{q+1}^{t+p} (B_{q+1}^{t+p})^T + (B_{q}^{t})^T (B_{q}^{t}).
$$

Although the expressions of the weighted persistent Laplacian matrices are different from the original persistent Laplacian matrices, some properties of $\mathcal{L}_{q}^{t+p}$ are preserved. The weighted persistent Laplacian operators are still symmetric and positive semi-defined. Additionally, their ranks are the same as $\mathcal{L}_{q}^{t+p}$. With the embedded volume information, weighted PSGs can provide richer topological and geometric information through the associated persistent Betti numbers and non-harmonic spectra (i.e., non-zero eigenvalues).

In real applications, we are more interested in the $0$, $1$, $2$-combinatorial Laplacian matrices because it is more intuitive to depict the relation among vertex, edges, and faces. Given a set of vertices $V = \{v_0, v_2, \cdots, v_N\}$ with $N + 1$ isolated points (0-simplices) randomly arranged in $\mathbb{R}^n$. By varying the radius $r$ of the $(n - 1)$-sphere centered at each vertex, a variety of simplicial complexes is created. We denote the simplicial complex generated at radius $r$ to be $K_r$, then the $0$-persistent $q$-combinatorial Laplacian operator and matrix at initial set up $K_r$ is

$$
\mathcal{L}_{q}^{0} = B_{q+1}^{0} (B_{q+1}^{0})^T + (B_{q}^{0})^T (B_{q}^{0}).
$$

The volume of any 1-simplex $\sigma_1 = [v_i, v_j]$ is $\text{Vol}(\sigma_1)$ and is actually the distance between $v_i$ and $v_j$, denoted $d_{ij}$. Then the $0$-persistent 0-combinatorial Laplacian matrix based on filtration $r$ can be expressed explicitly as

$$
(\mathcal{L}_{q}^{0})_{ij}^{r+0} = \begin{cases} 
-\sum_{j} (\mathcal{L}_{q}^{0})_{ij}^{r+0}, & \text{if } i = j \\
-1, & \text{if } i \neq j \text{ and } d_{ij} - 2r < 0 \\
0, & \text{otherwise}. 
\end{cases}
$$

Correspondingly, we can denote the 0-persistent 1-combinatorial Laplacian matrix based on filtration $r$ by $\mathcal{L}_{1}^{0}$, and the 0-persistent 2-combinatorial Laplacian matrix based on filtration $r$ by $\mathcal{L}_{2}^{0}$.

Alternatively, variants of persistent 0-combinatorial Laplacian matrices can be defined by adding the Euclidean distance information. The distance-weight persistent 0-combinatorial Laplacian matrix based on filtration $r$ can be expressed explicitly as

$$
(\mathcal{L}_{q}^{0})_{ij}^{r+0} = \begin{cases} 
-\sum_{j} (\mathcal{L}_{q}^{0})_{ij}^{r+0}, & \text{if } i = j \\
-d_{ij}, & \text{if } i \neq j \text{ and } d_{ij} - 2r < 0 \\
0, & \text{otherwise}. 
\end{cases}
$$

Moreover, the inverse-distance-weight persistent 0-combinatorial Laplacian matrix based on filtration $r$ can also be implemented:
The bond length of benzene is 1.39 Å, and the C–H bond length is around 1.08 Å, which are the real bond lengths in benzene molecule. Figure 5c shows that a 1-dimensional hole (1-cycle) is born when the filtration parameter \( r \) increases to 0.70 Å and dead when \( r = 1.21 \) Å. In Figure 5B,C, it can be seen that variants of 0-persistent 0-combinatorial Laplacian and 1-combinatorial Laplacian matrices based on filtration give us the identical \( \beta_0^{+0} \) and \( \beta_1^{+0} \) information, respectively.

Finally, it is mentioned that using the present procedure, more general weights, such as the radial basis function of ANM, to create a family of spectral graphs with different characteristic length scales for a given dataset. Similarly, the spectra of these matrices, the topological invariants and geometric shapes can be revealed from the given input point-cloud data.

2.3.5 | Multiscale spectral analysis

In the past few years, we have developed a multiscale spectral graph method such as generalized GNM and generalized ANM, to create a family of spectral graphs with different characteristic length scales for a given dataset. Similarly, in our PST, we can construct a family of spectral graphs induced by a filtration parameter. Moreover, we can sum over all the multiscale spectral graphs as an accumulated spectral graph. Specifically, a family of spectral graphs can be generated via the filtration. By analyzing the persistent spectra of these matrices, the topological invariants and geometric shapes can be revealed from the given input point-cloud data.

The spectra of \( L_{0}^{r+0}, L_{0}^{r+0} \), and \( L_{0}^{r+0} \) mentioned above carry similar information on how the topological structures of a graph are changed during the filtration. Benzene molecule (C\(_6\)H\(_6\)), a typical aromatic hydrocarbon, which is composed of six carbon atoms bonded in a planar regular hexagon ring with one hydrogen joined with each carbon atom. It provides a good example to demonstrate the proposed PST. Figure 4 illustrates the filtration of the benzene molecule. Here, we label 6 hydrogen atoms by H\(_1\), H\(_2\), H\(_3\), H\(_4\), H\(_5\), and H\(_6\), and the carbon adjacent to the labeled hydrogen atoms connect and constitute a single component, which gives rise to \( \beta_0^{+0} = 1 \). Furthermore, we can deduce that the C–C bond length of the benzene ring is about 1.40 Å, and the C–H bond length is around 1.08 Å, which are the real bond lengths in benzene molecule. Figure 5c shows that a 1-dimensional hole (1-cycle) is born when the filtration parameter \( r \) increases to 0.70 Å and dead when \( r = 1.21 \) Å. In Figure 5B,C, it can be seen that variants of 0-persistent 0-combinatorial Laplacian and 1-combinatorial Laplacian matrices based on filtration give us the identical \( \beta_0^{+0} \) and \( \beta_1^{+0} \) information, respectively.

The C–C bond length of benzene is 1.39 Å, and the C–H bond length is 1.09 Å. Due to the perfect hexagon structure of the benzene ring, we can calculate all of the distances between atoms. The shortest and longest distances...
between carbons and the hydrogen atoms are 1.09 Å and 3.87 Å. In Figure 5A, a total of 10 changes of $\lambda_{r=0}^{+}$ occur at various radii. Table 4 lists all the distances between atoms and the values of radii when the changes of $\lambda_{r=0}^{+}$ occur. It can be seen that the distance between atoms approximately equals twice of the radius value when a
jump of \((\bar{\lambda}_2)_{0}^{r+0}\) occurs. Therefore, we can detect all the possible distances between atoms with the nonzero spectral information. Moreover, in Figure 5B, the values of the smallest nonzero eigenvalues of \(L_{0}^{r+0}, \bar{L}_{0}^{r+0}\), and \(\bar{L}_{0}^{r+0}\) change concurrently.

3 | APPLICATIONS

In this section, we apply the proposed PST to the study of two important systems, fullerenes and proteins. All three different types of persistent combinatorial Laplacian matrices are employed in our investigation. The resulting persistent spectra contain not only the full set of topological persistence from the harmonic spectra, which is identical to that from a persistent homology analysis, but also non-harmonic eigenvalues and eigenvectors. Since the power of topological persistence has been fully explored and exploited in the past decade,\textsuperscript{20-23} to demonstrate the additional utility of our persistent spectral analysis, we mainly emphasize the non-harmonic spectra in the present applications. In particular, we demonstrate that PST is able to accurately predict the B-factors of a set of proteins for which the current biophysical models break down.

3.1 | Fullerene analysis and prediction

In 1985, Kroto et al discovered the first structure of C\textsubscript{60},\textsuperscript{41} which was confirmed by Kratschmer et al in 1990.\textsuperscript{42} Since then, the quantitative analysis of fullerene molecules has become an interesting research topic. The understanding of the fullerene structure–function relationship is important for nanoscience and nanotechnology. Fullerene molecules are only made of carbon atoms that have various topological shapes, such as the hollow spheres, ellipsoids, tubes, or rings. Due to the monotony of the atom type and the variety of geometric shapes, the minor heterogeneity of fullerene structures can be ignored. The fullerene system offers a moderately large dataset with relatively simple structures. Therefore, it is suitable for validating new computational methods because every single change in the spectra is interpretable. The proposed PST, that is, persistent spectral analysis, is applied to characterize fullerene structures and predict their stability.

All the structural data can be downloaded from CCL.NET Webpage. This dataset gives the coordinates of fullerene carbon atoms. In this section, we will analyze fullerene structures and predict the heat of formation energy.

3.1.1 | Fullerene structure analysis

The smallest member of the fullerene family is C\textsubscript{20} molecule with a dodecahedral cage structure. Note that 12 pentagons are required to form a closed fullerene structure. Following the Euler’s formula, the number of vertices, edges, and faces on a polygon have the relationship \(V - E + F = 2\). Therefore, the 20 carbon atoms in the dodecahedral cage form 30 bonds with the same bond length. The C\textsubscript{20} is the only fullerene smaller than C\textsubscript{60} that has the molecular symmetry of the full icosahedral point group \(I_h\). C\textsubscript{60} is a molecule that consists of 60 carbon atoms arranged as 12 pentagon rings and 20 hexagon rings. Unlike C\textsubscript{20}, C\textsubscript{60} has two types of bonds: 6 : 6 bonds and 6 : 5 bonds. The 6 : 6 bonds are shorter than 6 : 5 bonds, which can also be considered as “double bond.”\textsuperscript{43} C\textsubscript{60} is the most well-known fullerene with geometric symmetry \(I_h\). Since C\textsubscript{20} and C\textsubscript{60} are highly symmetrical, they are ideal systems for illustrating the persistent spectral analysis.

Figure 6A illustrates the radius filtration process built on C\textsubscript{20}. As the radius increases, the solid balls corresponding to carbon atoms grow, and a sequence of \(L_{0}^{r+0}\) matrices can be defined through the overlap relations among the set of balls. At the initial state \((r = 0.00 \text{ Å})\), all of the atoms are isolated from one another. Therefore, \(L_{0}^{r+0}\) is a zero matrix with dimension 20 \(\times\) 20. Since the C\textsubscript{20} molecule has the same bond length, which can be denoted as \(l(C_{20})\), once the radius of solid balls is greater than \(l(C_{20})\), all of the balls are overlapped, which makes the system a singly connected component. Figure 6B depicts the accumulated \(L_{0}^{r+0}\) for C\textsubscript{20}. For C\textsubscript{60}, the accumulated \(L_{0}^{r+0}\) is described in Figure 7A. Figure 7B-F shows the plots of \(L_{0}^{r+0}\) under different filtration \(r\) values. The blue cell located at the \(i\)th row and \(j\)th columns means the balls centered at atom \(i\) and atom \(j\) were connected to each other, that is, a 1-simplex formed with its vertex was \(i\) and \(j\). When the radius filtration increases, more and more bluer cells are created. In Figure 7F, the color of cells, except the cells located in the diagonal, turns to blue, which means all of the carbon atoms are connected with one another at \(r = 3.6\ \text{Å}\). For clarity, we set the diagonal terms to 0.
In Figure 8, the blue solid line represents $C_{20}$ properties and the dash orange line represents $C_{60}$ properties. For Figure 8A, the blue line drops at $r = 0.72$ Å, which means that the bond length of $C_{20}$ is around 1.44 Å. The orange line drops at $r = 0.68$ Å and 0.72 Å, which means that the “double bond” length of $C_{60}$ is around 1.36 Å and the 6 : 5 bond length is around 1.44 Å. Moreover, the total number of “double bond” is 30, yielding $\beta^0_{r + 0} = 30$ when the radius of solid balls is over 0.68 Å. In conclusion, one can deduce the number of different types of bonds as well as the bond length information from the number of zero eigenvalues (ie, $\beta^0_{r + 0}$) under the radius filtration. Furthermore, the geometric
information can also be derived from the plot of \( \lambda_2^{r+0} \). Each jump in Figure 8D at a specific radius represents the change of geometric and topological structures. The smallest non-zero eigenvalue \( \lambda_2^{r+0} \) of matrices for \( C_{20} \) changes 5 times in Figure 8D, which means \( C_{20} \) has five different distances between carbon atoms. Furthermore, by Remark 2.1, as \( \lambda_2^{r+0} \) of \( C_{20} \) keeps increasing, the smallest vertex connectivity of the connected subgraph continues growing and the topological structure becomes steady. As can be seen in the right-corner chart of Figure 6, the carbon atoms will finally grow to a solid object with a steady topological structure.

Figure 8B depicts the changes of Betti 1 value \( \beta_1^{r+0} \) (ie, the number of zero eigenvalues for \( L_1^{r+0} \)) under the filtration \( r \). Since \( C_{20} \) has 12 pentagonal rings, \( \beta_1^{r+0} \) jumps to 11 when radius \( r \) equals to the half of the bond length of \( l(C_{20}) \). These eleven 1-cycles disappear at \( r = 1.17 \) Å. There are 12 pentagons and 20 hexagons in \( C_{60} \), which results in \( \beta_1^{r+0} = 12 \) at \( r = 0.72 \) Å and \( \beta_1^{r+0} = 31 \) at \( r = 1.17 \) Å. All of the pentagons and hexagons disappear at \( r = 1.22 \) Å.
As the filtration process, even more structure information can be derived from the number of zero eigenvalues of \( L_{r}^{\delta \sigma} + 0 \) (ie, \( \beta_{r}^{\delta \sigma} + 0 \)) in Figure 8C. For \( C_{20} \), \( \beta_{r}^{\delta \sigma} = 1 \) when \( r = 1.17 \) Å, which corresponds to the void structure in the center of the dodecahedral cage. The void disappears at \( r = 1.65 \) Å since a solid structure is generated at this point. For fullerene \( C_{60} \), 20 hexagonal cavities and a center void exist from 1.12 Å to 1.40 Å yielding \( \beta_{r}^{\delta \sigma} = 21 \). As the filtration goes, hexagonal cavities disappear which results \( \beta_{r}^{\delta \sigma} \) decrease to 1. The central void keeps alive until a solid block is formed at \( r = 3.03 \) Å. In a nutshell, we can deduce the number of different types of bonds, the bond length, and the topological invariants from the present persistent spectral analysis.

3.1.2 | Persistent spectral predictions of fullerene stability using both harmonic and non-harmonic spectra

Having shown that the detailed fullerene structural information can be extracted into the spectra of \( L_{r}^{\delta \sigma} + 0 \), we further illustrate that fullerene functions can be predicted from their structures by using our PST in this section. Similar structure-function analysis has been carried out by using other methods.\(^{24,33,44}\) For small fullerene molecule series \( C_{20} \) to \( C_{60} \), with the increase in the number of atoms, the ground-state heat of formation energies decrease.\(^{45,46}\) The left chart in Figure 9 describes this phenomenon. Similar patterns can also be found in the total energy (STO-3G/SCF at MM3) per atom and the average binding energy of \( C_{2n} \). To analyze these patterns, many theories have been proposed. Isolated pentagon rule assumes that the most stable fullerene molecules are those in which all the pentagons are isolated. Zhang et al.\(^ {46} \) stated that fullerene stability is related to the ratio between the number of pentagons and the number of carbon atoms. Xia and Wei\(^ {44} \) proposed that the stability of fullerene depends on the average number of hexagons per atom. However, these theories all focused on the pentagon and hexagon information. More specifically, they use purely topological information to reveal the stability of fullerene. In contrast, we believe that both harmonic persistent spectra and non-harmonic persistent spectra can be used to model the structure-function relationship of fullerenes. We hypothesize that the non-harmonic persistent spectra of \( L_{0}^{\delta \sigma} + 0 \) matrices are powerful enough to model the stability of fullerene molecules. To verify our hypothesis, we compute the summation, mean, maximal, SD, variance of its eigenvalues, and \( \langle \tilde{\lambda}_{j} \rangle_{r}^{\delta \sigma} + 0 \) of the persistent spectra of \( L_{0}^{\delta \sigma} + 0 \) over various filtration radii \( r \). We depict a plot with the horizontal axis representing radius \( r \) and the vertical axis representing the particular spectrum value, which is actually the same as Figure 8. Then we define the area under the plot of spectra with a negative sign as

\[
A_{\alpha} = -\sum_{i=1}^{\Lambda_{i}^{\alpha}} \delta r,
\]

where \( \delta r \) is the radius grid spacing, in Figure 8, \( \delta r = 0.01 \) Å. Here, \( \alpha = \text{Sum, Avg, Max, SD, Var, and Sec} \) being the type index, and thus, \( \Lambda_{i}^{\alpha} \) represent the summation, mean, maximal, SD, variance, and the smallest non-zero eigenvalue.
\( (\tilde{\lambda}_2)_0^{r+0} \) of \( L_0^{r+0} \) at \( i \)th radius step, respectively. The right chart in Figure 9 describes the area under the plot of spectra and closely resembles that of the heat of formation energy. We can see that generally the left chart and the middle chart show the same pattern. The integration of \( (\tilde{\lambda}_2)_0^{r+0} \) decreases as the number of carbon atoms increases. However, the structural data we used might not be the same ground-state data as in Reference 46, which results in \( C_{36} \) being not matching the corresponding energy perfectly. Limited by the availability of the ground-state structural data, we are not able to analyze the full set of the fullerene family.

As the present persistent spectral method provides both topological (ie, harmonic) and non-topological spectral analyses, we are interested in a further comparison of their performances. For the topological spectral analysis, we adopted the model of Xia and Wei,\(^{44}\) which was based on the atom number averaged Betti-2 length for the cavity. The Betti-2 length for the cavity is the longest persistence of \( \beta_2^{r+0} \) (ie, the number of zero eigenvalues for \( L_2^{r+0} \) under the filtration \( r \)). Table 5 presents the predictions obtained from both topological spectral analysis and non-topological spectral analysis. The right chart of Figure 9 presents the correlations between predicted energies and the heat of formation energy of the fullerene molecules computed from quantum mechanics.\(^{46}\) Clearly, both methods provide very accurate predictions.

To quantitatively validate our model, we apply one of the simplest machine learning algorithms, linear least-squares method, to predict the heat of formation energy. The Pearson correlation coefficient for the \( \alpha \)th method is defined as

\[
C^\alpha_c = \left\{ \sum_{k=1}^{N} \left( A_k^\alpha - \bar{A} \right) \left( E_k - \bar{E} \right) \right\} \left\{ \sum_{k=1}^{N} \left( A_k^\alpha - \bar{A} \right)^2 \sum_{k=1}^{N} \left( E_k - \bar{E} \right)^2 \right\}^{-\frac{1}{2}}
\]

where \( A_k^\alpha \) represents the theoretically predicted energy of the \( k \)th fullerene molecule, \( E_k \) represents the heat of formation energy of the \( k \)th fullerene molecule, and \( \bar{A} \) and \( \bar{E} \) are the corresponding mean values.

Table 6 lists the correlation coefficient under different type index \( \alpha \) and harmonic spectral information of \( L_2^{r+0} \). The highest correlation coefficient is close to unity (0.986) obtained with \( \alpha = \text{Max} \). The lowest correlation coefficient is 0.942 with \( \alpha = \text{Sum} \). We can see that all the correlation coefficients are close to unity, which verifies our hypothesis that the non-harmonic spectra of \( L_0^{r+0} \) have the capacity of modeling the stability of fullerene molecules. The performance of the topological spectra is on a bar with that of the best of the non-harmonic spectra. Obviously, our persistent harmonic spectral information and persistent non-harmonic spectral information could be trivially combined in the present machine learning setting to achieve an even higher accuracy. Therefore, our PST works extremely well only with both harmonic and non-harmonic spectra, which means our PST is a powerful tool for quantitative data analysis and prediction.

| Table 5 | The heat of formation energy of fullerenes\(^{46}\) and its corresponding energies predicted non-harmonic spectral model (\( \alpha = \text{Max} \)) and harmonic spectral model |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Fullerene type | \( C_{20} \) | \( C_{24} \) | \( C_{26} \) | \( C_{30} \) | \( C_{32} \) | \( C_{36} \) | \( C_{50} \) | \( C_{60} \) |
| Heat of formation energy | 1.180 | 1.050 | 0.989 | 0.850 | 0.781 | 0.706 | 0.509 | 0.401 |
| Non-harmonic spectral energy | 1.138 | 1.050 | 0.964 | 0.821 | 0.857 | 0.766 | 0.474 | 0.391 |
| Harmonic spectral energy | 1.224 | 0.962 | 0.929 | 0.861 | 0.825 | 0.737 | 0.564 | 0.364 |
| Note: The unit is EV/atom. |

| Table 6 | The correlation coefficients for predictions using different type of statistics of non-harmonic spectra and topological spectra (top) |
| --- | --- | --- | --- | --- | --- | --- | --- |
| Type index | Sum | Avg | Max | SD | Var | Sec | Top |
| Correlation coefficient | 0.942 | 0.985 | 0.986 | 0.969 | 0.977 | 0.981 | 0.979 |
3.2  |  Protein flexibility analysis

As clarified earlier, the number of zero eigenvalues of \( p \)-persistent \( q \)-Laplacian matrix (\( p \)-persistent \( q \)th Betti number) can also be derived from persistent homology. Persistent homology has been used to model fullerene stability.\(^{44}\) In this section, we further illustrate the applicability of present PST by a case that non-harmonic persistent spectra offer a unique theoretical model, whereas it may be difficult to come up with a suitable persistent homology model for this problem. To this end, we consider a challenging biophysical problem for which current methods do not work well.

The protein flexibility is known to correlate with a wide variety of protein functions. It can be modeled by the beta factors or B-factors, which are also called Debye-Waller factors. B-factors are a measure of the atomic mean-square displacement or uncertainty in the X-ray scattering structure determination. Therefore, understanding the protein structure, flexibility, and function via the accurate protein B-factor prediction is a vital task in computational biophysics.\(^{47}\) Over the past few years, quite many methods are developed to predict protein B-factors, such as GNM,\(^{30}\) ANM,\(^{31}\) FRI,\(^{48,49}\) and MWCG.\(^{34,47}\) However, all of the aforementioned methods are based on a particular matrix derived from the graph network, which is constructed using alpha carbon as nodes and connections between nodes as edges. In this section, we apply our PST to create richer geometric information in B-factor prediction. In particular, we are interested in the prediction of B-factors for which the standard GNM fails to work.

To illustrate our method, we consider a protein whose total number of residues is \( N \). In this work, we employ the coarse-grained \( C_\alpha \) representation. Similarly, like in the previous application of fullerene structure analysis, we treat each \( C_\alpha \) atom as a 0-simplex at the initial setup and assign it a solid ball with a radius of \( r \). By varying the filtration parameter \( r \), we can obtain a family of \( L_0^{r+0} \). For each matrix \( L_0^{r+0} \), its corresponding ordered spectrum is given by

\[
(\lambda_1)^{r+0}_0, (\lambda_2)^{r+0}_0, \ldots, (\lambda_N)^{r+0}_0.
\]

Suppose the number of zero eigenvalues is \( m \), then we have \( \beta_0^{r+0} = m \). Since \( L_0^{r+0} \) is symmetric, then eigenvectors of \( L_0^{r+0} \) corresponding to different eigenvalues must be orthogonal to each other. The Moore-Penrose inverse of \( L_0^{r+0} \) can be calculated by the non-harmonic spectra of \( L_0^{r+0} \):

\[
(L_0^{r+0})^{-1} = \sum_{k=m+1}^{N} \frac{1}{(\lambda_k)^{r+0}_0} (u_k)^{r+0}_0 (u_k)^{r+0}_0^T,
\]

where \( T \) is the transpose and \( (u_k)^{r+0}_0 \) is the \( k \)th eigenvector of \( L_0^{r+0} \). The modeling of \( i \)th B-factor at filtration parameter \( r \) can be expressed as

\[
B_i^r = (L_0^{r+0})^{-1}, \forall i = 1, 2, \ldots, N,
\]

**FIGURE 10**  Comparison of protein B-factors obtained by experiment (left chart), PST (middle chart), and GNM (right chart) for protein PDB ID: 1V70, visualized in Visual Molecular Dynamics (VMD).\(^{51}\) Red color represents for the most flexible regions.
and the final model of $i$th B-factor is given by

$$B_{i}^{PST} = \sum_{r} w_r B_{r} + w_0, \forall i = 1, 2, \cdots, N,$$

TABLE 7  Pearson correlation coefficients in B-factor predictions using GNM, NMA, and PST for a group of proteins that are very challenging to GNM

<table>
<thead>
<tr>
<th>PDB ID</th>
<th>GNM</th>
<th>NMA</th>
<th>PST</th>
</tr>
</thead>
<tbody>
<tr>
<td>1OPD</td>
<td>0.398</td>
<td>0.372</td>
<td>0.886</td>
</tr>
<tr>
<td>2PKT</td>
<td>−0.286</td>
<td>−0.245</td>
<td>0.745</td>
</tr>
<tr>
<td>2FQ3</td>
<td>0.384</td>
<td>0.609</td>
<td>0.808</td>
</tr>
<tr>
<td>1R7J</td>
<td>0.368</td>
<td>0.400</td>
<td>0.908</td>
</tr>
<tr>
<td>1W2L</td>
<td>0.397</td>
<td>0.130</td>
<td>0.808</td>
</tr>
<tr>
<td>5CYT</td>
<td>0.331</td>
<td>0.248</td>
<td>0.624</td>
</tr>
<tr>
<td>1V70</td>
<td>0.162</td>
<td>0.265</td>
<td>0.826</td>
</tr>
<tr>
<td>1CCR</td>
<td>0.351</td>
<td>0.379</td>
<td>0.708</td>
</tr>
<tr>
<td>2VIM</td>
<td>0.212</td>
<td>0.243</td>
<td>0.535</td>
</tr>
<tr>
<td>2CG7</td>
<td>0.379</td>
<td>0.355</td>
<td>0.780</td>
</tr>
<tr>
<td>1QTO</td>
<td>0.334</td>
<td>0.672</td>
<td>0.853</td>
</tr>
<tr>
<td>1WHI</td>
<td>0.270</td>
<td>0.492</td>
<td>0.612</td>
</tr>
<tr>
<td>1NKO</td>
<td>0.368</td>
<td>0.583</td>
<td>0.770</td>
</tr>
<tr>
<td>1O08</td>
<td>0.309</td>
<td>0.281</td>
<td>0.811</td>
</tr>
<tr>
<td>2HQK</td>
<td>0.348</td>
<td>0.715</td>
<td>0.808</td>
</tr>
</tbody>
</table>

Note: Pearson correlation coefficients for GNM and NMA are extracted from Reference 50.
where \( w \) and \( w_0 \) are fitting parameters which can be determined from a simple machine learning algorithm, the linear regression, using B-factors from experimental data \( B_{\text{Exp.}} \). In this application, we consider the filtration radius from 2 to 6 with the grid spacing of 0.5 and grid spacing of 2 from 6 to 26, then totally 20 different \( L_{r=0}^r \) are created at \( r = 1.5, 2, 2.5, 3, 3.5, 4, 4.5, 5, 5.5, 6, 8, 10, 12, 14, 16, 18, 20, 22, 24, \) and 26. By calculating all the non-harmonic spectra together with their eigenvectors, 20 Moore-Penrose inverse matrices \( (L_{r=0}^r)^{-1} \) are constructed.

Park et al have carried out the B-factor prediction of small-, medium-, and large-sized proteins using GNM and NMA. We collected a total of 15 medium-sized or large-sized proteins for which the Pearson correlation coefficient of GNM prediction is less than 0.4 reported in an earlier study. Figure 10 plots the flexibility of 1V70 with red color representing for the most flexible regions (the highest B-factor). Obviously, B-factors predicted by PST are very similar to experimental values, whereas GNM predictions differ dramatically from experimental B-factors. This situation can also be seen from Figure 11 where B-factors predicted by PST and GNM are compared with those of experimental results. GNM fails to work for 1V70, 1R7J, and 2FQ3. The Pearson correlation coefficients of GNM, NMA, and PST are compared in Table 7. This study shows that the proposed PST has a great potential in multiscale biophysical modeling and prediction.

4 CONCLUSION

SGT is a powerful tool for data analysis due to its ability to extract geometric and topological information. However, its performance can be quite limited for various reasons. One of them is that the current SGT does not provide a multiscale analysis. Motivated by persistent homology and multiscale graphs, we introduce PST as a unified paradigm to unveil both topological persistence and geometric shape from high-dimensional datasets.

For a point set \( V \subset \mathbb{R}^n \) without additional structures, we construct a filtration using an \((n-1)\) sphere of a varying radius \( r \) centered at each point. A series of persistent combinatorial Laplacian matrices are induced by the filtration. It is noted that our harmonic persistent spectra (ie, zero eigenvalues) fully recover the persistent barcode or persistent diagram of persistent homology. Specifically, the numbers of zero eigenvalues of persistent \( q \)-combinatorial Laplacian matrices are the \( q \)-dimensional persistent Betti numbers for the same given filtration. However, additional valuable spectral information is generated from the non-harmonic persistent spectra. In this work, in addition to persistent Betti numbers and the smallest non-zero eigenvalues, five statistic values, namely, sum, mean, maximum, SD, and variance, are also constructed for data analysis. We use a few simple two-dimensional (2D) and three-dimensional (3D) structures to carry out the proof-of-principle analysis of the PST. The detailed structural information could be incorporated into the persistent spectra. For instance, for the benzene molecule, the approximate C—C bond and C—H bond length can be intuitively read from the plot of the 0-dimensional persistent Betti numbers. Moreover, PST also has the capacity to accurately predict the heat of formation energy of small fullerene molecules. We use the area under the plot of the persistent spectra to model fullerene stability and apply the linear least-squares method to fit our prediction with the heat of formation energy. The resulting correlation coefficient is close to 1, which shows that our PST has an excellent performance on molecular data. Furthermore, we have applied our PST to the protein B-factor prediction. In this case, persistent homology does not offer a straightforward model. We consider a set of 15 challenging proteins for which the most popular biophysical method, Gaussian network model (GNM), fails to work. We show that the additional non-harmonic persistent spectral information provides extremely successful B-factor predictions to this set of challenging proteins.

It is pointed out that the proposed persistent spectral analysis can be paired with advanced machine learning algorithms, including various deep learning methods, for a wide variety of applications in data science. In particular, the further construction of element-specific PST and its application to protein-ligand binding affinity prediction and computer-aided drug design will be reported elsewhere.

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CONFLICT OF INTEREST

The author declare no conflict of interest.
DATA AVAILABILITY STATEMENT
Data are available at https://weilab.math.msu.edu/PSG/.

ENDNOTES
1If \( q = 0, \partial_q \) is a zero map, and we denote \( \partial_0 \) a zero matrix with dimension \( 1 \times N \), where \( N \) is the number of 0-simplices. Note that this term is needed to attain the correct dimension of the null space. If \( q > \text{dim}(K) \), we will not discuss \( \partial_q \) since there is no \( \sigma_q \) in \( K \).
2We define the boundary matrix \( B_q \) for boundary map \( \partial_q \) as a zero matrix. The number of columns of \( B_0 \) is the number of 0-simplices in \( K \), the number of rows will be 1.
3We carry out feature scaling to make sure all \( B_q \) are on a similar scale.

REFERENCES


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