Intrinsic Sliced Wasserstein Distances for Comparing Collections of Probability Distributions on Manifolds and Graphs

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Abstract
Collections of probability distributions arise in a variety of applications ranging from user activity pattern analysis to brain connectomics. In practice these distributions can be defined over diverse domain types including finite intervals, circles, cylinders, spheres, other manifolds, and graphs. This paper introduces an approach for detecting differences between two collections of distributions over such general domains. To this end, we propose the intrinsic slicing construction that yields a novel class of Wasserstein distances on manifolds and graphs. These distances are Hilbert embeddable, allowing us to reduce the distribution collection comparison problem to a more familiar mean testing problem in a Hilbert space. We provide two testing procedures one based on resampling and another on combining p-values from coordinate-wise tests. Our experiments in various synthetic and real data settings show that the resulting tests are powerful and the p-values are well-calibrated.

1. Introduction

Distributional data defined over general domains such as manifolds and graphs arise in a variety of statistical applications. In this paper we consider the problem of comparing two collections of distributions over such a general domain. Our goal is to test for homogeneity—whether all of the distributions come from the same meta-distribution—in an interpretable manner. While conceptually similar to two-sample testing, this is a higher order notion in the sense that our units of analysis are distributions/histograms.

For instance, given collections of personal activity histograms (over cylinder: time of day \times intensity) for two sub-populations, one may be interested in determining whether there are statistically significant differences between activity patterns of these sub-populations. As another example, consider normalized counts of events per geographic region on a daily basis. Collected over a year, this gives a set of 365 daily probability distributions over the region adjacency graph, and one may wish to compare the collection of distributions from weekdays to those from weekends. Testing specific aspects of homogeneity is preferable: for example, in regular two-sample testing, detecting that the means are unequal provides interpretable insights, whereas a general test that only says there are unspecified differences between the distributions is less useful for interpretation.

Limited settings of this problem tackling distributions over the interval/circle have been considered in the literature (Dubey & Müller, 2019), yet the general case of distributions over graphs and manifolds is open. The requirement to test for specific differences is non-trivial on general domains: what is the equivalent of mean for a collection of distributions? While Fréchet mean (Peyré & Cuturi, 2019) may seem like the natural choice, there are a number of problems with testing Fréchet mean equality. First, the existence and uniqueness of the Fréchet mean is not guaranteed, and it can be sensitive to small changes. Second, computing the Fréchet mean is expensive and can become prohibitive when resampling is used to compute the null distribution. Finally, resampling poses conceptual problems: using permutation null will detect differences beyond the equality of Fréchet means (same problem exists in regular two-sample testing, see e.g. Huang et al. (2006)), and using bootstrap requires designing the null case, which is highly non-trivial.

We attack this problem using insights from recent developments that utilize Hilbert embeddings for simplifying distributional data problems (Solomon et al., 2014; Petersen & Müller, 2016). The simplification comes as a result of linearity of Hilbert spaces, which allows adapting existing statistical approaches to distributional data. A crucial requirement on the embedding is that the distance in the embedding space should give a meaningful distance between measures; it is this property that renders quantities computed in the embedding space such as means and variances meaningful. While transportation based distances are effi-
We overcome these difficulties by introducing a new slicing whereby rejections lead to interpretable insights. We provide constructions suitable for manifolds or graphs. The Sliced Wasserstein (SW) distance and its generalized variant (Kolouri et al., 2019a, GSW) sets up the idea of approximating Wasserstein distances using multiple nonlinear projections, it is presented in extrinsic terms (i.e. Euclidean space) and can suffer from the curse of dimensionality when a low dimensional data manifold lives in a high-dimensional space. Our choice of eigenfunctions for projection is very different from the one-parameter function families in GSW and allows us to rigorously prove a number of general and testing-specific properties.

Moreover, the GSW construction does not directly apply to graphs. While the tree-sliced variant of GSW (Le et al., 2019) can be applied in an intrinsic manner (the clustering variant), it relies on a different type of distance, in the limit related to the euclidean/geodesic distance. This can be seen by comparing our lower bound to theirs: our lower bound for ISW is in terms of the MMD using the spectral distance (Proposition 4.4). The recently-proposed Sobolev transport (Le et al., 2022) does consider measures supported on graphs, but in absence of a slicing construction it is limited to testing for unspecific differences and requires extensive compute due to much slower permutation based calibration (see Section 6). Deshpande et al. (2019) proposed Max Sliced Wasserstein (MSW) distance—taking maximum over projected $W_2$ distances vs the average $W_2$ distances of SW—in the context of generative models.

Finally, the robust sliced Wasserstein distance of Lai & Zhao (2017) does make use of the geometric properties of the underlying manifold. However, their goal is to compute a correspondence between two manifolds by mapping them into $\mathbb{R}^d$ using eigenmaps and treating the mapped manifolds as measures in $\mathbb{R}^d$ and minimizing some version of Euclidean slicing. None of the aforementioned works consider the problem of comparing collections of distributions, provide ways for obtaining calibrated $p$-values, or prove properties of the distances that make them desirable for hypothesis testing.

\section{2. Related Work}

Our framework is not simply a higher order version of a two-sample kernel test (Gretton et al., 2012) since we test for equality of a specific aspect of meta-distributions. This renders our null hypothesis different from Gretton et al. (2012), and we need a different set of techniques both for proofs and computations. For example, testing in Gretton et al. (2012) can use the permutation null which is valid due to the stronger null hypothesis of equal distributions. In contrast, with our null hypothesis we cannot use the permutation null and have to resort to a bootstrap procedure. Other approaches such as the general Hilbert embedding framework of Petersen & Müller (2016) is not tied to a distance between probability distributions and so can be problematic for capturing the location and variability aspects of distribution collections. In addition, Petersen & Müller (2016) has difficulties in higher dimensions and does not provide constructions suitable for manifolds or graphs.
3. Preliminaries

3.1. Problem Setup

Given a compact metric space $\mathcal{X}$, let $\mathcal{P}(\mathcal{X})$ denote the set of Borel probability measures on $\mathcal{X}$. Our main interest is in the case where $\mathcal{X}$ is a graph or a manifold with the shortest/geodesic distance as the metric, and thus the compactness restriction. The 2-Wasserstein distance can be defined on $\mathcal{P}(\mathcal{X})$ using the metric of $\mathcal{X}$ as the ground distance (Peyré & Cuturi, 2019; Panaretos & Zemel, 2019), giving $\mathcal{W}_2^2(\mathcal{X}) : \mathcal{P}(\mathcal{X}) \times \mathcal{P}(\mathcal{X}) \to \mathbb{R}_{\geq 0}$. Due to the repeated use of the real line case we use the shorthand $\mathcal{W}_2$ when $\mathcal{X} = \mathbb{R}$, i.e., $\mathcal{W}_2 := \mathcal{W}_2^2$. Central to our study are distributions on the space of probability measures $\mathcal{P}(\mathcal{P}(\mathcal{X})) = (\mathcal{P}(\mathcal{X}), B(\mathcal{P}(\mathcal{X})))$, where $B(\mathcal{P}(\mathcal{X}))$ is the Borel $\sigma$-algebra generated by the topology induced by $\mathcal{W}_2^2(\mathcal{X})$ (Bigot, 2020). To avoid confusion, we will refer to the elements of $\mathcal{P}(\mathcal{P}(\mathcal{X}))$ as meta-distributions.

Let $P, Q \in \mathcal{P}(\mathcal{P}(\mathcal{X}))$, and assume that we are given two collections of probability measures $\{\mu_i\}_{i=1}^{N_1}$ and $\{\nu_i\}_{i=1}^{N_2}$ that are drawn from $P$ and $Q$: $\mu_i \sim P$ and $\nu_i \sim Q$ in an independent-and-identically-distributed (i.i.d.) manner. Our goal is to use this sample to test the null hypothesis of whether $P = Q$. While this is conceptually a two-sample test, note that our data points are distributions; in practice, the distributions $\mu_i$ or $\nu_i$ are given by histograms.

**Remark 3.1.** Let us compare this with the usual two-sample testing. Consider $P \in \mathcal{P}(\mathcal{P}(\mathcal{X}))$ constructed as follows. Let $\mu^* \in \mathcal{P}(\mathcal{X})$ be a fixed probability measure. Let $x_1, x_2, \ldots, x_A \sim \mu^*$ and construct the histogram summarizing this sample: $\frac{1}{A} \sum_{a=1}^{A} \delta_{x_a}$. Now, $\frac{1}{A} \sum_{a=1}^{A} \delta_{x_a} \in \mathcal{P}(\mathcal{X})$ is one sample drawn from $P$. Suppose one gets the collection $\{\mu_i\}_{i=1}^{N_1}$, where each histogram is obtained as above: $\mu_i \sim \mu^*$. Similarly, consider $Q \in \mathcal{P}(\mathcal{P}(\mathcal{X}))$ of the same type based on some other fixed $\nu^* \in \mathcal{P}(\mathcal{X})$, and let $\{\nu_i\}_{i=1}^{N_2}$ the corresponding collection of histograms. Testing whether $P = Q$ in the limit boils down to $\mu^* = \nu^*$. When compared to the usual two-sample testing this may seem rather inefficient, requiring $A$ times more samples (resp. $N_1A$ and $N_2A$ samples from $\mu^*$ and $\nu^*$). However, in our general setup it is **not assumed** that the histograms in the collections come from meta-distributions of the above simple type (i.e. all $\mu_i$ are generated by drawing from the same underlying distribution $\mu^*$). In fact, the target use-case for our approach is when these histograms are collected by observing different individuals who have their **person-specific** behaviors/distributions.

**Remark 3.2.** To gain further insight into our problem setup, consider it through the lens of statistical manifold theory (Murray & Rice, 1993, SMT). Assume that we observe two collections of parametric distributions: $\mu_i = F(x, \beta_i), \beta_i \sim G(\beta)$ for $i = 1, \ldots, N_1$, and $\nu_i = F(x, \gamma_i), \gamma_i \sim H(\gamma)$ for $i = 1, \ldots, N_2$. Thus, each collection is generated from a distribution of its parameters, resp. $G(\cdot)$ and $H(\cdot)$. The goal of our test is to find out whether $G$ and $H$ are the same solely by observing the collections $\{\mu_i\}_{i=1}^{N_1}$ and $\{\nu_i\}_{i=1}^{N_2}$. While SMT and the Fisher information metric provides a powerful framework for studying parametric families of probability distributions, we focus on more general spaces of probability measures and the Fisher metric may not be easily extended to such nonparametric spaces of measures.

3.2. Hilbert Embeddings

Let $D(\cdot, \cdot) : \mathcal{P}(\mathcal{X}) \times \mathcal{P}(\mathcal{X}) \to \mathbb{R}_{\geq 0}$ be a distance between probability distributions. $D(\cdot, \cdot)$ is called **Hilbertian** (this is just a naming convention; no implication that the map is a Hilbert map) if there exist a Hilbert space $\mathcal{H}$ and a map $\eta : \mathcal{P}(\mathcal{X}) \to \mathcal{H}$ such that $D(\mu, \nu) = \|\eta(\mu) - \eta(\nu)\|_\mathcal{H}$. For example, it is well-known that 2-Wasserstein distance $\mathcal{W}_2^2(\mathcal{X})$ on general $\mathcal{X}$ is not Hilbertian (Peyré & Cuturi, 2019); however, the 2-Wasserstein distance $\mathcal{W}_2^2$ on general $\mathcal{X}$ is Hilbertian (Peyré & Cuturi, 2019).

Since the map $\eta$ takes every measure on $\mathcal{X}$ to a point in $\mathcal{H}$, we see that a meta-distribution $P \in \mathcal{P}(\mathcal{P}(\mathcal{X}))$ gives a rise to a measure on $\mathcal{H}$ given by pushforward operation, $\eta\#P = P \circ \eta^{-1} \in \mathcal{P}(\mathcal{H})$. In addition, if a finite dimensional approximation $\eta_D : \mathcal{P}(\mathcal{X}) \to \mathbb{R}^D$ of $\eta$ is available, then $\eta_D\#P$ is a measure on $\mathbb{R}^D$. This observation is enormously useful: problems about the elements of the rather abstract space $\mathcal{P}(\mathcal{P}(\mathcal{X}))$ are reduced to problems about familiar measures on $\mathcal{H}$ or even $\mathbb{R}^D$. For example, the usual notions of mean and variance can be applied to the measure $\eta\#P$ to gain insights about the meta-distribution $P$. The validity of these insights hinges on the $\eta$-map coming from a Hilbertian distance, as distances are central to the statistical quantities of interest.

Testing for $\eta\#P = \eta\#Q$ can serve as a proxy for our original testing problem of $P = Q$. As typical with two-sample tests, various aspects of the equality $\eta\#P = \eta\#Q$ can be tested, such as the mean or variance equality; unspecified tests of equality can be applied as well. We will concentrate on testing certain aspects of the equality so that one can easily drill down on the results. This is similar to the regular two-sample testing where checking for equality of, say, means is often preferable as it gives immediately interpretable insights, whereas a general test that only says there are unspecified differences between the distributions is less useful for interpretation. To obtain succint and interpretable tests we concentrate on the mean of the resulting pushforward measure in $\mathcal{H}$.

**Definition 3.3.** For a meta-distribution $P \in \mathcal{P}(\mathcal{P}(\mathcal{X}))$, define its **Hilbert centroid** with respect to the Hilbertian
distance $D$ as $C_{\eta \# P} := E_{\mu \sim P}[\eta(\mu)] \in H$, assuming it exists.

Our testing procedure is based on checking the equality $C_{\eta \# P} = C_{\eta \# Q}$, or more explicitly: $E_{\mu \sim P}[\eta(\mu)] = E_{\nu \sim Q}[\eta(\nu)]$. Intuitively, each “dimension” of the map $\eta$ probes some aspect of the two involved meta-distributions and makes sure that they are in agreement in expectation. One of our testing approaches will use the statistic

$$T(P, Q) := ||C_{\eta \# P} - C_{\eta \# Q}||_H^2.$$  (1)

to capture the deviations from equality; this quantity can be written directly in terms of pairwise distances.

**Proposition 3.4.** For $P, Q \in P(P(X))$, the following holds:

$$T(P, Q) = E_{\mu \sim P, \nu \sim Q}[D^2(\mu, \nu)] - \frac{1}{2}E_{\mu, \nu \sim P}[D^2(\mu, \mu')] - \frac{1}{2}E_{\nu, \nu' \sim Q}[D^2(\nu, \nu')] .$$

Next we give an example of what Hilbert centroid equality implies in an important special case.

**Example 1.** Let $X = [0, T] \subset \mathbb{R}$ with $D$ being the 2-Wasserstein distance $W_2$. Given a probability measure $\mu \in P([0, T])$, let $F_\mu$ be its cumulative distribution function: $F_\mu(x) = \mu([0, x]) = \int_0^x d\mu$. The generalized inverse of cumulative distribution function (CDF) is defined by $F^{-1}_\mu(s) := \inf\{x \in [0, T] : F_\mu(x) > s\}$. The squared 2-Wasserstein distance has a rather simple expression in terms of the inverse CDF (Peyré & Cuturi, 2019):

$$W^2_2(\mu, \nu) = \int_0^1 (F^{-1}_\mu(s) - F^{-1}_\nu(s))^2 ds.$$  (2)

This formula immediately establishes the Hilbertianity of $W_2$ through the map $\eta : P([0, T]) \rightarrow L_2([0, T])$ defined by $\eta(\mu) = F^{-1}_\mu$. Note that $\eta$ is invertible for increasing normalized functions in the embedding space. Using this insight, we see that the corresponding “average measure” of $P \in P(P(X))$ can be introduced via $P_{av} = \eta^{-1}(E_{\mu \sim P}[\eta(\mu)])$. It is easy to prove that $P_{av}$ satisfies the following: $P_{av} = \arg\min_{P \in P(X)} E_{\mu \sim P}[W_2(\mu, \rho)^2]$, which is the definition of the Fréchet mean, see for example (Peyré & Cuturi, 2019). In this setting, $C_{\eta \# P} = C_{\eta \# Q}$ boils down to having the same Fréchet means, $P_{av} = Q_{av}$.

We will later see that the Hilbert embedding corresponding to the intrinsic sliced 2-Wasserstein distance is assembled of embeddings like in Example 1 applied after pushforwards (see Figure 1 for an intuition). This means that the resulting equality $C_{\eta \# P} = C_{\eta \# Q}$ becomes more stringent, making it a better proxy for detecting the deviations from $P = Q$ without losing the interpretability aspect.

### 4. Intrinsic Sliced 2-Wasserstein Distance

We introduce a Hilbertian version of $W_2$ on manifolds and graphs via a construction we call intrinsic slicing due to its use of the domain’s intrinsic geometric properties. To focus our discussion we concentrate on the manifold case, as the graph case is simpler and is obtained by replacing the Laplace-Beltrami operator by the graph Laplacian.

Let $\lambda_\ell, \phi_\ell : \ell = 0, 1, \ldots$ be the eigenvalues and eigenfunctions of the Laplace-Beltrami operator on $X$ with Neumann boundary conditions. The eigenfunctions are sorted by increasing eigenvalue and assumed to be orthonormal with respect to some fixed (e.g. uniform) measure on $X$; also $\phi_0 = const$ and $\lambda_0 = 0$. One can define the spectral kernel $k(x, y) = \sum_\ell \alpha(\lambda_\ell)\phi_\ell(x)\phi_\ell(y)$ and the corresponding spectral distance on the manifold $d(x, y) = k(x, x) + k(y, y) - 2k(x, y) = \sum_\ell \alpha(\lambda_\ell)(\phi_\ell(x) - \phi_\ell(y))^2$, where $\alpha : \mathbb{R}_+ \rightarrow \mathbb{R}_+$ is a function that controls contribution from each spectral band. By setting $\alpha(\lambda) = e^{-\lambda}$ for some $t > 0$, we get the heat/diffusion kernel and the corresponding diffusion distance (Coifman & Lafon, 2006).

Another important case is $\alpha(\lambda) = 1/\lambda^2$ if $\lambda > 0$ and $\alpha(0) = 0$, which gives the biharmonic kernel and distance (Lipman et al., 2010). In both of these constructions $\alpha(\cdot)$ is a decreasing function, allowing the smoother low-frequency (i.e. smaller $\lambda_\ell$) eigenfunctions to contribute more.

#### 4.1. Definition and properties

A real-valued function $\phi : X \rightarrow \mathbb{R}$ can be used to map the manifold $X$ onto the real line. Any probability measure $\mu \in P(X)$ can likewise be projected onto the real line using the pushforward of $\phi$, which we denote by $\phi_\# \mu = \mu \circ \phi^{-1} \in P(\mathbb{R})$. While the pushforward notions used here and in previous sections are conceptually the same, for clarity we use $\sharp$ for measures and $\#$ for meta-distributions.

We define intrinsic slicing as follows.

**Definition 4.1.** Given a function $\alpha : \mathbb{R}_+ \rightarrow \mathbb{R}_+$ and a probability distribution $D(\cdot, \cdot)$ on $P(\mathbb{R})$, we define the intrinsic sliced distance $ISD(\cdot, \cdot)$ on $P(X)$ by

$$ISD^2(\mu, \nu) = \sum_\ell \alpha(\lambda_\ell)D^2(\phi_\ell \# \mu, \phi_\ell \# \nu).$$

The choice of the Laplacian eigenfunctions in the definition can be justified by a number of their properties. Eigenfunctions are intrinsic quantities of a manifold and are ordered with respect to some fixed (e.g. uniform) measure on $X$; also $\phi_0 = const$ and $\lambda_0 = 0$. One can define the spectral kernel $k(x, y) = \sum_\ell \alpha(\lambda_\ell)\phi_\ell(x)\phi_\ell(y)$ and the corresponding spectral distance on the manifold $d(x, y) = k(x, x) + k(y, y) - 2k(x, y) = \sum_\ell \alpha(\lambda_\ell)(\phi_\ell(x) - \phi_\ell(y))^2$, where $\alpha : \mathbb{R}_+ \rightarrow \mathbb{R}_+$ is a function that controls contribution from each spectral band. By setting $\alpha(\lambda) = e^{-\lambda}$ for some $t > 0$, we get the heat/diffusion kernel and the corresponding diffusion distance (Coifman & Lafon, 2006).

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resulting distance \textit{Intrinsic Sliced 2-Wasserstein Distance}, and denote it by $\mathcal{IS}W_2$. First, we discuss the convergence of the infinite sum in Definition 4.1.

\textbf{Proposition 4.2.} If $\mathcal{X}$ is a smooth compact $n$-dimensional manifold and $\sum_\ell \lambda_\ell^{(n-1)/2} \alpha(\lambda_\ell) < \infty$, then $\mathcal{IS}W_2$ is well-defined.

Next, we prove a number of properties of $\mathcal{IS}D$.

\textbf{Proposition 4.3.} If $\mathcal{D}$ is a Hilbertian probability distance such that $\mathcal{IS}D$ is well-defined, then (i) $\mathcal{IS}D$ is Hilbertian, and (ii) $\mathcal{IS}D$ satisfies the following metric properties: non-negativity, symmetry, the triangle inequality, and $\mathcal{IS}D(\mu, \mu) = 0$.

\textbf{Proof.} By Hilbertian property of $\mathcal{D}$, there exists a Hilbert space $\mathcal{H}$ and a map $\eta^\mu : \mathcal{P}(\mathbb{R}) \to \mathcal{H}$ such that $\mathcal{D}(\mu_1, \mu_2) = \|\eta^\mu(\mu_1) - \eta^\mu(\mu_2)\|_\mathcal{H}$ for all $\mu_1, \mu_2 \in \mathcal{P}(\mathbb{R})$. Plugging this into Definition 4.1 we have $\mathcal{IS}D(\mu, \nu) = \|\eta(\mu) - \eta(\nu)\|_\mathcal{H}$, where $\mathcal{H} = \hat{e}_i \mathcal{H}$ and the $\ell$-th component of $\eta(\mu)$ is $\sqrt{\alpha(\lambda_\ell)} \eta(\phi_\ell \mu) \in \mathcal{H}$. The second part of Proposition 4.3 directly follows from the Hilbert property.

Since $\mathcal{W}_2$ is Hilbertian on $\mathcal{P}(\mathbb{R})$, the application of Proposition 4.3 yields that $\mathcal{IS}W_2$ is also Hilbertian, making it possible to use $\mathcal{IS}W_2$ for our hypothesis tests in Section 5.

For a simple choice of distance $\mathcal{D}$ on $\mathcal{P}(\mathbb{R})$, namely the absolute mean difference, the corresponding intrinsic sliced distance is the well-known MMD (Gretton et al., 2012).

\textbf{Proposition 4.4.} Let $\mathcal{D}(\mu_1, \mu_2) = |\mathbb{E}_{x \sim \mu_1} [x] - \mathbb{E}_{y \sim \mu_2} [y]|$ for $\mu_1, \mu_2 \in \mathcal{P}(\mathbb{R})$, then the corresponding $\mathcal{IS}D$ is equivalent to the MMD with the spectral kernel $k(\cdot, \cdot)$.

When $k(x, y)$ is the heat kernel, the sliced distance in Proposition 4.4 is very much like the MMD with the Gaussian kernel, with the parameter $t$ in $\alpha(\lambda) = e^{-t\lambda}$ controlling the kernel width. Indeed, the two kernels coincide on $\mathbb{R}$, and on general manifolds Varadhan’s formula gives asymptotic equivalence for small $t$ (Berger, 2003).

An interesting insight derived from the above result is that $\mathcal{IS}W_2$ is in a sense a “stronger” distance than MMD that uses the corresponding spectral kernel. The $\mathcal{IS}W_2$ compares the quantiles of the pushforward distributions (Eq. 2), whereas MMD compares their expectations only. We formalize this notion in the next result, also providing a theoretical reason for preferring $\mathcal{IS}W_2$ for hypothesis testing.

\textbf{Proposition 4.5.} $\text{MMD}(\mu, \nu) \leq \text{IS}W_2(\mu, \nu)$ when the same $\alpha(\cdot)$ is used in both constructions.

We are now in a position to prove that $\mathcal{IS}W_2$ is a true metric.

\textbf{Theorem 4.6.} If $\alpha(\lambda) > 0$ for all $\lambda > 0$, then $\mathcal{IS}W_2$ is a metric on $\mathcal{P}(\mathcal{X})$.

We remind that 2-Wasserstein distance can be defined directly on $\mathcal{P}(\mathcal{X})$ using the geodesic distance as the ground metric; we denote this distance as $\mathcal{W}_2^{X}$. Lipschitz properties of the eigenfunctions imply the following:

\textbf{Proposition 4.7.} There exists a constant $c$ depending only on $\mathcal{X} \subseteq \mathbb{R}^n$ such that for all $\mu, \nu \in \mathcal{P}(\mathcal{X})$ the inequality $\mathcal{IS}W_2(\mu, \nu) \leq c \mathcal{W}_2^X(\mu, \nu) \sqrt{\sum_\ell \lambda_\ell^{(n+3)/2} \alpha(\lambda_\ell)}$ holds.

Our final result looks at the quantity $\mathcal{T}$ defined using $\mathcal{IS}W_2$ by Eq. (1). We will be using $\mathcal{T}$ computed on finite collections of measures as a test statistic in the next section. We show that it enjoys robustness with respect to small perturbations of the measures in the collection.

\textbf{Proposition 4.8.} Let $\{\mu_i\}_{i=1}^N$ and $\{\nu_i\}_{i=1}^N$ be two collections of probability measures on $\mathcal{P}(\mathcal{X})$, such that $\|\nu_i - \mathcal{W}_2^X(\mu_i, \nu_i) \leq \epsilon$, then $\mathcal{T}(\{\mu_i\}_{i=1}^N, \{\nu_i\}_{i=1}^N) \leq C \epsilon^2$. Here $C = c \sqrt{\sum_\ell \lambda_\ell^{(n+3)/2} \alpha(\lambda_\ell)}$ from previous proposition and is assumed to be finite.

This bound implies that if the distributions in a collection undergo horizontal shifts that are small as measured by the geodesic distance $\mathcal{W}_2^X$, then $\mathcal{T}$ is small as well.

4.2. Approximate Hilbert Embedding

An important aspect of $\mathcal{IS}W_2$ is that its Hilbert map $\eta : \mathcal{P}(\mathcal{X}) \to \mathcal{H}$ can be approximated by a finite-dimensional embedding $\eta_D : \mathcal{P}(\mathcal{X}) \to \mathbb{R}^D$ such that $\mathcal{IS}W_2^D(\mu, \nu) \approx \|\eta_D(\mu) - \eta_D(\nu)\|_{\mathbb{R}^D}$. This is useful for practical computation and for one of our hypothesis testing approaches.

Using the formula for $\mathcal{IS}W_2$ on $\mathcal{P}(\mathbb{R})$ in terms of the quantile function, Eq. (2), the Hilbert map is defined by $\eta^\mu(\lambda) = F^{-1}_{\mu}(\lambda)$. We have $\mathcal{W}_2^D(\mu, \nu) \approx \|\eta^\mu(\lambda) - \eta^\nu(\lambda)\|_{\mathbb{R}^D}$, where the norm involves integration. We can discretize the integral using the Riemann sum for equidistant knots $s_k = k/\ell$, $k = 1, ..., \ell$, define the approximate embedding $\eta_D^0 : \mathcal{P}(\mathbb{R}) \to \mathbb{R}^D$ as:

$$\eta_D^0 : \mu \to \frac{1}{\sqrt{\ell}} \left[ F^{-1}_{\mu}(s_1), ..., F^{-1}_{\mu}(s_\ell) \right]. \hspace{1cm} (3)$$

Now, $\mathcal{W}_2(\mu, \nu) \approx \|\eta_D^0(\mu) - \eta_D^0(\nu)\|_{\mathbb{R}^D}$ with approximation quality depending on the embedding dimension $\ell$.

To approximate the Hilbert map for $\mathcal{IS}W_2$ we truncate the series defining $\mathcal{IS}W_2$ and use a finite number of eigenfunctions for pushforward: $\phi_\ell, \ell = 1, ..., \ell$, where $\phi_0$ is dropped since it is a constant. By inspecting the proof of Proposition 4.3 and using Eq. (3), we can define $\eta_D : \mathcal{P}(\mathcal{X}) \to \mathbb{R}^D$ with $D = \ell D'$ as the concatenation of $\ell$ maps:

$$(\eta_D)\ell : \mu \to \frac{\alpha(\lambda)}{\ell} \left[ F^{-1}_{\phi_\ell \mu}(s_1), ..., F^{-1}_{\phi_\ell \mu}(s_{\ell}) \right].$$

Spectral decompositions of Laplace-Beltrami operators for general manifolds (Coifman & Lafon, 2006; Reuter et al.,...}

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We show in Appendix A.4 that a) summands in the second family can be computed analytically (see Appendix A.3).

The hyperparameters $L$ and $D'$ capture distinct aspects of the complexity of the distribution. A larger $L$ allows access to higher order eigenfunctions that possess greater spatial oscillations, linking to the finer details of the geometry and topology of the underlying domain and distribution collection. On the other hand, $D'$ captures how well the pushforward distribution is represented, which is determined by the number of quantiles. Both hyperparameters can have a significant impact on the success of the testing process.

5. Hypothesis Testing

Let $\{\mu_i\}_{i=1}^{N_1}$ and $\{\nu_i\}_{i=1}^{N_2}$ be two i.i.d. collections of measures drawn from $P, Q \in \mathcal{P}(\mathcal{P}(X))$ respectively. Our goal is to use these samples to test the null hypothesis $H_0 : C_{\eta#P} = C_{\eta#Q}$, where $\eta$ is the Hilbert embedding of the sliced distance $ISW_2$ on $\mathcal{P}(X)$.

5.1. Resampling Based Test

We use the quantity $T(\cdot, \cdot)$ from Eq. (1) as the test statistic. Its sample version is computed by replacing the expectations by the empirical means, and excluding the diagonal terms to achieve unbiasedness

$$\tilde{T} = \sum_{i,j: i \neq j} ISW_2^2(\mu_i, \nu_j) / 2N_1(N_1 - 1) + \sum_{i,j: i \neq j} ISW_2^2(\nu_i, \mu_j) / 2N_2(N_2 - 1)$$

(4)

$$- \sum_{i,j} ISW_2^2(\mu_i, \nu_j) / N_1 N_2.$$

Note that $E\tilde{T} = T(P, Q)$. In practice, the $ISW_2$ values are computed from the approximate embedding: $ISW_2(\mu_1, \mu_2) \approx \|\eta_D(\mu_1) - \eta_D(\mu_2)\|_{\mathcal{D}}$. We denote the resulting statistic by $\tilde{T}_{L,D'}$.

The difference between $\tilde{T}_{L,D'}$ and the population version (i.e. $T(\cdot, \cdot)$ can be decomposed as $(\tilde{T} - T) + (T - T_L) + (T_L - \tilde{T}_{L,D'})$, where the sums inside the terms $\tilde{T}_L$ and $T_{L,D'}$ correspond to partial sums that approximate $ISW_2^2(\cdot, \cdot)$ by $\sum_{i=1}^L \alpha(\lambda_i) W^2_D(\phi_i, \phi_i)$, and $W^2_D(\phi_i, \phi_i')$ by $\|\eta_D(\phi_i) - \eta_D(\phi_i')\|^2$, respectively. We show in Appendix A.4 that a) summands in the second and third terms in the sum can be made infinitesimally small by choosing large enough $L$ and $D'$, respectively; b) an asymptotic result for the first difference can be obtained by extending the tools from Gretton et al. (2012) and Serfling (2009). These results are based on several assumptions detailed in Appendix A.4. Combining the two results, we establish asymptotic distributions of $\tilde{T}_{L,D'}$.

**Theorem 5.1.** Assume relevant conditions (see Appendix A.4) hold. Define $N = N_1 + N_2$, and suppose that as $N_1, N_2 \to \infty$, we have $N_1/N \to \rho_1, N_2/N \to \rho_2 = 1 - \rho_1$, for some fixed $0 < \rho_1 < 1$. With $L \geq L_N, D' \geq D_N$ chosen in an appropriate way (see Appendix A.4), under $H_0 : C_{\eta#P} = C_{\eta#Q}$ we have

$$N \tilde{T}_{L,D'} \sim \sum_{m=1}^{\infty} \gamma_m(A_m^2 - 1),$$

where $A_m \sim N(0, 1)$ for $m = 1, 2, \ldots$, and $\gamma_m$ are the eigenvalues of a certain operator that depends on $P$ and $Q$.

Further, under $H_1 : C_{\eta#P} \neq C_{\eta#Q}$, $\sqrt{N} (\tilde{T}_{L,D'} - 1)$ is asymptotically Gaussian with mean $0$ and finite variance.

We evaluate the power performance of the testing procedure based on $\tilde{T}_{L,D'}$ for the sequence of contiguous alternatives $H_{1N} = \{(P, Q) : C_{\eta#P} = C_{\eta#Q} + \delta_N, l = 1, 2, \ldots\}$, where the deviation from null is quantified collectively by pushforward differences $\delta_N \in \mathcal{H}$, $\delta_N = \oplus_i \sqrt{\langle \gamma_i \delta_N \rangle}$ that are made to approach 0 as $N \to \infty$. The following theorem establishes consistency of our testing procedure against a family of such local alternatives.

**Theorem 5.2.** Assume the same conditions and choice of $L, D'$ as Theorem 5.1. Then for the sequence of contiguous alternatives $H_{1N}$ such that $N\|\delta_N\|_{\mathcal{D}}^2 \to \infty$, the test based on $\tilde{T}_{L,D'}$ is consistent for any $\alpha \in (0, 1)$, that is as $N \to \infty$ the asymptotic power approaches 1.

**Testing Procedure** In practice, to obtain the $p$-value for the $\tilde{T}_{L,D'}$-statistic we use a bootstrap procedure. Remember that $\tilde{T}_{L,D'}$ is computed via the approximate embedding $\eta_D$ with $D = LD'$. The collection $\{\mu_i\}_{i=1}^{N_1}$ is mapped to the collection $\{X_i = \eta_D(\mu_i)\}_{i=1}^{N_1}$ of vectors in $\mathbb{R}^D$ drawn in an i.i.d. manner from $\eta_D#P = P \circ \eta_D^{-1} \in \mathcal{P}(\mathbb{R}^D)$. Similarly, for the other collection we have a sample $\{Y_i = \eta_D(\nu_i)\}_{i=1}^{N_2}$ drawn from $\eta_D#Q$. Now, the null $H_0 : C_{\eta#P} = C_{\eta#Q}$ implies that the means of the distributions $\eta_D#P$ and $\eta_D#Q$ coincide in $\mathbb{R}^D$.

The bootstrap null distribution for $\tilde{T}_{L,D'}$ can be obtained as follows. Let $X$ and $Y$ be the sample means; construct the combined sample $\{X_i = X + \frac{X_i - X}{\sqrt{N_1}}\}_{i=1}^{N_1} \cup \{Y_i = Y + \frac{Y_i - Y}{\sqrt{N_2}}\}_{i=1}^{N_2}$. This centers both samples at $\frac{X+Y}{2}$. Now, from the combined sample we select with replacement $N_1$ (resp. $N_2$) samples to make bootstrap sample $\{X_i^{N_1}\}_{i=1}^{N_1}$ (resp. $\{Y_i^{N_2}\}_{i=1}^{N_2}$). Repeat this process $B$ times (we take $B = 1000$ in our experiments), and collect the null test statistic values $\tilde{T}_{L,D'}^B = \tilde{T}_{L,D'}(\{X_i^{N_1}\}_{i=1}^{N_1}, \{Y_i^{N_2}\}_{i=1}^{N_2})$ for $b = 1, \ldots, B$. The approximate $p$-value is then given by:

$$p = \frac{1}{B+1} \left( \sum_{b=1}^B \mathbb{1}(\tilde{T}_{L,D'} \geq \tilde{T}_{L,D'}) + 1 \right).$$

**Remark 5.3.** Permutation testing cannot be applied here as it would detect differences beyond the mean inequality.
Such differences help reject the stronger null hypothesis $H_0 : P = Q$ which can be the intent in some situations. However, such a rejection will not allow pinpointing the aspect responsible for the difference; see (Huang et al., 2006).

5.2. Testing via $p$-value Combination

The bootstrap test above incurs a high computational cost and the granularity of the $p$-values is determined by the number of resamples, which can be too coarse in massive multiple comparison settings often seen in industrial applications. Thus, we propose an approach that avoids resampling.

As explained above, testing $H_0 : C_{\eta\#}p = C_{\eta\#}Q$ can be interpreted as testing whether the means of the distributions $\eta_D\#P$ and $\eta_D\#Q$ coincide in $\mathbb{R}^D$. To this end, we adopt the approach proposed by Rustamov & Klosowski (2020) in a spatial statistics context.

First, we apply the Behrens-Fisher-Welch $t$-test (Liu & Xie, 2020), but in our numerical experiments we found that the Cauchy combination approach encounters problems when any of the $p$-values is very close to 1, which can happen in our setting due to the form of the embedding $\eta_D$. Therefore, in contrast to Rustamov & Klosowski (2020), for us the harmonic combination is the only appropriate choice.

To guarantee size control, we establish a version of Theorem 1 from Liu & Xie (2020) for the harmonic mean $p$-value. Assume that a test statistic $Z \in \mathbb{R}^D$ has null distribution with zero mean and every pair of coordinates of $Z$ follows bivariate Gaussian distribution. Compute the coordinate-wise two-sided $p$-values $p_k$, $k = 1, 2, ..., D$. Second, an overall $p$-value is computed via the harmonic mean $p$-value combination method which is robust to dependencies (Good, 1958; Wilson, 2019):

$$p^H = H \left( D / \left( \frac{1}{p_1} + \frac{1}{p_2} + \cdots + \frac{1}{p_D} \right) \right),$$

where the function $H$ has a known form described in (Wilson, 2019).

Another approach for combining $p$-values is the Cauchy combination test (Liu & Xie, 2020), but in our numerical experiments we found that the Cauchy combination approach encounters problems when any of the $p$-values is very close to 1, which can happen in our setting due to the form of the embedding $\eta_D$. Therefore, in contrast to Rustamov & Klosowski (2020), for us the harmonic combination is the only appropriate choice.

To guarantee size control, we establish a version of Theorem 1 from Liu & Xie (2020) for the harmonic mean $p$-value. Assume that a test statistic $Z \in \mathbb{R}^D$ has null distribution with zero mean and every pair of coordinates of $Z$ follows bivariate Gaussian distribution. Compute the coordinate-wise two-sided $p$-values $p_k$, $k = 1, 2, ..., D$. Second, an overall $p$-value is computed via the harmonic mean $p$-value combination method which is robust to dependencies (Good, 1958; Wilson, 2019):

$$p^H = H \left( D / \left( \frac{1}{p_1} + \frac{1}{p_2} + \cdots + \frac{1}{p_D} \right) \right),$$

where the function $H$ has a known form described in (Wilson, 2019).

6. Experiments

We compare the performance of our tests with several existing methods, across synthetic and real data, and settings of the embedding parameters $L, D'$. For evaluation, we use empirical power at different degrees of departure from the null hypothesis (Captured by $\delta$ in the plots in Fig. 3), calculated by averaging the proportion of rejections at level $\alpha = 0.05$ over 1000 independent datasets, with samples divided into two groups of sizes $n_1 = 60, n_2 = 40$. To ensure the tests are well-calibrated, we calculate nominal sizes assuming the two sample groups are drawn from the same meta-distribution. Details on the experiment settings and computational complexity are in Appendix B.1.

Finite intervals We use embedding dimensions $L = 3, D' = 10$ to compare our method against 11 functional ANOVA tests—we report results for 3 of them in Figure 3a (see Appendix B for complete results). All methods maintain nominal size for $\delta = 0$ (Figure 3a). While the combination test (ISD comb) based on our proposal outperformed all the other tests across all values of $\delta$, the bootstrap test that uses the overall T statistic (ISD T boot) performs better than Fmaxb but worse than others. In terms of computation time, ISD comb takes the least amount of time (Fig. 2a).

We also compare the $p$-value combination test based on an unsliced 24-dimensional inverse CDF embedding with sliced ISW2-based tests (Figure 3b). We use multiple pairs of $(L, D')$ values, all of them giving overall embeddings of dimension $D = LD' = 24$. The performance of an ISW2-based test that uses slicing over only the first eigenfunction is almost as good as the unsliced version. With more eigenfunctions, the powers first improve considerably, then become similar to the unsliced version again.

Manifold domains We consider data from distributions on circles and cylinders. For circular data, we take von Mises distributions with randomly chosen parameters as our samples. For an angle $x$ (measured in radians), the von Mises probability density function is given by $f(x|\mu, \kappa) = \exp[\kappa \cos(x - \mu)](2\pi I_0(\kappa))^{-1}$, where $I_0(\kappa)$ is the modified Bessel function of order 0. We fix $\kappa = 2$, and use $\mu \equiv \mu_i \sim N(0, 0.1^2)$, $\mu \equiv \nu_i \sim N(\delta, 0.1^2)$ for samples
Intrinsic Sliced Wasserstein Distances on Manifolds and Graphs

Figure 3. Performances on synthetic data. Dotted lines indicates nominal size of all tests ($\alpha = 0.05$). (a) comparison with existing methods on finite interval—a test based on basis function representation (Gorecki & Smaga, 2015, FP), a sum-type $\ell_2$ norm-based test (L2b) (Zhang, 2013), and a max-type test (Zhang et al., 2019) that uses the maximum of coordinate-wise $F$ statistic ($F_{maxb}$), (b) unsliced vs. different settings of $(L, D')$ on finite interval. (c) Circular data, comparing with Fréchet ANOVA (Dubey & Müller, 2019), and the DISCO nonparametric test (Rizzo & Székely, 2010); (d) harmonic combination tests on cylindrical data for $L = 4$.

from group 1 and 2 respectively—with $\delta \in [0, 15] \times \pi/180$ (i.e. 0 to 15 degrees converted to radians). As each observation vector, we take 100 random draws from each sample-specific distribution. For our embeddings, we use $L = 10, D' = 20$. Since the competing methods cannot handle circular geometry directly, to implement them we cut the circle into an interval. Figure 3c shows that all methods maintain nominal size, but both our tests maintain considerably higher power than existing methods for all $\delta$. Computationally, ISD comb has comparable order of magnitude as the other two methods (Fig. 2b).

To generate cylindrical data, we use the distribution proposed by Mardia & Sutton (1978). Samples from each distribution have the form of a bivariate random vector $(\Theta, X)$, where the first (circular) marginal $\Theta$ has a von Mises distribution, and $X$ is a Gaussian conditional on $\Theta = \theta$. We draw the mean parameters for each coordinate-wise distribution from $\text{Unif}(0, 1)$ and $\text{Unif}(\delta, \delta + 1)$ for sample groups 1 and 2 respectively, with $\delta \in [0, 30] \times \pi/180$. We then use 500 random draws from each distribution to obtain histograms. To evaluate the effects of choosing $L, D'$ we calculate our embeddings for $L \in \{2, 3, 4, 5\}, D' \in \{6, 8, 12, 24, 48\}$. The choice of $L$ has small effect on performance, so we report results for $L = 4$ in Figure 3d. Higher values of $D'$ result in some increase in power.

**Comparison to existing slicing methods** As an example showcasing the importance of using intrinsic distances, consider the following graph setup: points A and B are adjacent on a planar regular 200-gon, with the edge AB removed. The intrinsic distance between A and B is large and the (extrinsic) Euclidean distance is small. No matter what Euclidean projection we use, the SW cost of moving probability mass from A to B would be small, leading to testing power loss when we compare two distribution sets that concentrate one around A and the other around B. To numerically analyze this case we generate distributions on this graph by putting $\text{Unif}(0,1)$ weights at each of the vertices, and added an additional weight of 10 to the bin at point A to obtain the first set of distributions with a mode at A. For the second set we put the weight of 10 at B instead to obtain a set of distributions with mode at B.

Table 1 shows comparison of $\text{ISW}_2$ with Sliced Wasserstein (SW) in two dimensions, GSW with 3 choices of defining functions (circular, homogeneous polynomial of degree 3, and degree 5), MSW, and Sobolev transport (ST). We use $L = 10, D' = 8$ to produce all embeddings, and implement the sliced version of ST by aggregating ST distances originating from 10 random vertices. $\text{ISW}_2$ achieves the highest possible power, maintains nominal type-I error, and has computational time comparable to SW/GSW. ST and MSW achieve the same power as $\text{ISW}_2$, but take much longer. This is because unlike slicing-based methods, they need to rely on permutation testing in absence of embeddings. For MSW, the high power comes at the price of a size that is much higher than the acceptable threshold of 0.05.

**7. Real Data Examples**

**NHANES** This dataset (NHANES, 2008) contains physical activity pattern readings for 6839 individuals, corresponding to activity monitor intensity values for $24 \times 60 =$
We summarize the p-values for the actual data comparisons. Above diagonal: null p-values obtained by combining and randomly splitting the two groups. Bold entries correspond to rejected hypotheses with the BH procedure at FDR level 0.1.

**Table 2.** Activity intensity comparison across age groups in the NHANES data. Below diagonal: p-values for the actual data comparisons. Above diagonal: null p-values obtained by combining and randomly splitting the two groups. Bold entries correspond to rejected hypotheses with the BH procedure at FDR level 0.1.

| Age Groups | 6–15 | 16–25 | 26–35 | 36–45 | 46–55 | 56–65 | 66–75 | 76–85 |
|------------|------|-------|-------|-------|-------|-------|-------|-------|
| 6–15       | 0.394| 0.098 | 0.555 | 0.882 | 0.985 | 0.919 | 0.997 |
| 16–25      | 1.2e-13| 0.575 | 0.967 | 0.126 | 0.921 | 0.911 | 0.977 |
| 26–35      | 3.1e-21| 2.7e-04| 0.459 | 0.197 | 0.996 | 0.919 | 0.565 |
| 36–45      | 6.1e-22| 7.9e-08| 0.042 | 0.864 | 0.637 | 0.849 | 0.991 |
| 46–55      | 8.2e-22| 4.7e-05| 0.011 | 0.343 | 0.841 | 0.165 | 0.554 |
| 56–65      | 1.3e-25| 0.001 | 0.001 | 5.6e-05| 0.003 | 0.991 | 0.962 |
| 66–75      | 3.6e-35| 7.8e-12| 1.5e-11| 4.6e-15| 1.8e-13| 0.001 | 0.989 |
| 76–85      | 3.8e-46| 1.4e-26| 1.7e-30| 8.4e-37| 2.1e-35| 1.3e-17| 6.5e-09|

**Table 3.** Chicago Crime analysis p-values. Bold entries correspond to rejected hypotheses with the BH procedure at FDR level 0.1.

| Crime Type       | Tue vs Thu | Tue vs Sat |
|------------------|------------|------------|
| Theft            | 0.428      | **4.2e-06**|
| Deceptive Pract. | 0.313      | **0.001**  |
| Battery          | 0.430      | **0.001**  |
| Robbery          | 0.119      | **0.003**  |
| Narcotics        | 0.854      | **0.004**  |
| Criminal Dam.    | 0.855      | **0.02**   |
| Other Offense    | 0.931      | 0.052      |
| Burglary         | 0.142      | 0.261      |
| Assault          | 0.997      | 0.38       |
| Motor Veh. Theft | 0.858      | 0.416      |

1440 minutes throughout the day, over 7 days. We capture this activity pattern into a cylindrical histogram with time and intensity dimensions, having 96 and 100 bins respectively. Since the time dimension is periodic, normalized counts of this histogram can be considered as person-specific probability distributions over the cylinder $S^1(T_1) \times [0, T_2)$, with $T_1 = 96, T_2 = 100$. To check if activity patterns vary across different groups of individuals, we first split individuals into age-specific groups: 6–15, 16–25, ... 76–85, then sample 100 males and 100 females from each split. For our analysis, we consider $L = 3$ indices along the two directions, i.e. $\ell_1, \ell_2 = 1, 2, 3$ and $D' = 5$. We summarize the p-value combination test results in Table 6, **below the diagonal**. We control false discovery rate at 0.1 by running the resulting p-values through the procedure of Benjamini & Hochberg (1995). $ISW_2$ detects statistically significant differences between all pairs of groups, except the 36–45 and 46–55 groups. As expected, the control p-values—obtained by mixing male and female samples in each age group and splitting arbitrarily—do not concentrate near zero. More details are given in Appendix B.2.

**Chicago Crime** We use this dataset (City of Chicago, 2022) to show a practical usage of our method on histograms over graphs. Each beat (geographic area subdivision used by police) corresponds to a vertex, and two vertices are connected by an edge if the corresponding beats share a geographic boundary. For each crime type and day, the normalized counts of that crime type for each beat gives a daily probability distribution over the graph. Our goal is to compare the collection of distributions of, say, theft occurring on Tuesday to those of Thursday and Saturday. The Tuesday versus Thursday comparison is intended as a null case, as we do not expect to see any differences between them (Rustamov & Klosowski, 2020). Table 7 results $ISW_2$ outcomes using 100-dimensional embeddings ($L = 20, D' = 5$). We detect statistically significant differences between Tuesday and Saturday patterns for six categories of crime, and as expected, no differences between Tuesday and Thursday patterns. For more details and another graph-based application, see Appendices B.3 and B.4, respectively.

**8. Conclusion**

In this paper we have introduced a novel class of sliced Wasserstein distances on manifolds and graphs, and applied the resulting $ISW_2$ distance in the context of hypothesis tests for meta-distributions on general domains, due to the high relevance of this setup in real data situations. It is worth pointing out that the assumption-lean nature of $ISW_2$ can enable its adoption by many other statistical and ML methods.

The $ISW_2$ embeddings can be used to do other hypothesis tests, such as tests for equality of covariances by extending the notion of Wasserstein covariances from (Petersen & Müller, 2019) to general domains. The embeddings can also be used as input features for supervised learning problems over distributions. In another direction, $ISW_2$ is directly applicable as a loss function for generative modeling. For example, if one is generating distributions over a graph or manifold, $ISW_2$ can serve as a loss function similarly to how other sliced distances are used in this context over Euclidean domains.

The proposed framework has a few limitations. There is scope of exploration for choosing the parameters $L$ and $D'$ in a principled manner. Empirical computation of eigenfunctions for general manifolds will introduce approximation errors that need to be tackled by expanding our theoretical results. In terms of potential societal impacts of our proposal, any difference between data distributions from different demographic groups found using our method should be evaluated in light of potential biases in the data collection stage.
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Appendix

For the appendix to be self-contained, we restate results (theory and experiments) in the main paper when necessary. Appendix A contains proofs of theoretical results and some implementation details. Details and results of experiments performed are in Appendix B. Code and data behind these experiments are at https://github.com/shubhobm/isw.

A. Proofs and additional results

A.1. Proofs and Notes for Section 3

Proposition 3.4. For \( P, Q \in \mathcal{P}(\mathcal{P}(X)) \), the following equality holds:

\[
T(P, Q) = \mathbb{E}_{\mu \sim P, \nu \sim Q}[\mathcal{D}^2(\mu, \nu)] - \frac{1}{2} \mathbb{E}_{\mu, \mu' \sim P}[\mathcal{D}^2(\mu, \mu')] - \frac{1}{2} \mathbb{E}_{\nu, \nu' \sim Q}[\mathcal{D}^2(\nu, \nu')],
\]

where to avoid notational clutter we use \( \mathcal{D}^2(\cdot, \cdot) \) as a shorthand for \( (\mathcal{D}(\cdot, \cdot))^2 \).

Proof. This is a straightforward application of the “kernel trick”: using the Hilbert property of the distance we can rewrite,

\[
\mathbb{E}_{\mu \sim P, \nu \sim Q}[\|\eta(\mu) - \eta(\nu)\|_H^2] - \frac{1}{2} \mathbb{E}_{\mu, \mu' \sim P}[\|\eta(\mu) - \eta(\mu')\|_H^2] - \frac{1}{2} \mathbb{E}_{\nu, \nu' \sim Q}[\|\eta(\nu) - \eta(\nu')\|_H^2]
\]

\[
= \mathbb{E}_{\mu \sim P}[\|\eta(\mu)\|_H^2] + \mathbb{E}_{\nu \sim Q}[\|\eta(\nu)\|_H^2] - 2(\mathbb{E}_{\mu \sim P}[\eta(\mu)], \mathbb{E}_{\nu \sim Q}[\eta(\nu)])_H
\]

\[
- \mathbb{E}_{\mu, \mu' \sim P}[\|\eta(\mu)\|_H^2] + \mathbb{E}_{\nu \sim Q}[\|\eta(\nu)\|_H^2]
\]

\[
+ (\mathbb{E}_{\mu \sim P}[\eta(\mu)], \mathbb{E}_{\nu \sim Q}[\eta(\nu)])_H + (\mathbb{E}_{\nu \sim Q}[\eta(\nu)], \mathbb{E}_{\nu \sim Q}[\eta(\nu)])_H
\]

\[
= \|\mathbb{E}_{\mu \sim P}[\eta(\mu)] - \mathbb{E}_{\nu \sim Q}[\eta(\nu)]\|_H^2 = T(P, Q).
\]

Which gives the sought equivalence.

A.2. Proofs and Notes for Section 4.1

Proposition 4.2. If \( X \) is a smooth compact \( n \)-dimensional manifold and \( \sum \lambda_\ell^{(n-1)/2} \alpha(\lambda_\ell) < \infty \), then \( ISW_2 \) is well-defined.

Proof. We use Hörmander’s bound on the supremum norm of the eigenfunctions:

\[
\|\phi_\ell\|_\infty \leq c\lambda_\ell^{(n-1)/4}\|\phi_\ell\|_2,
\]

for some constant \( c \) that depends on the manifold. By orthonormality of the eigenfunctions we have \( \forall \ell, \|\phi_\ell\|_2 = 1 \). Next, note that \( \forall \ell, \|\phi_\ell\|_\infty \leq 2\|\phi_\ell\|_\infty \) as the maximum distance that the mass would be transported in any transportation plan involving pushforwards via \( \phi_\ell \) is upper bounded by \( 2\|\phi_\ell\|_\infty \). As a result, every term in the series defining \( ISW_2 \) can be upper-bounded by the terms of the following series:

\[
\sum \lambda_\ell^{(n-1)/2} \alpha(\lambda_\ell)
\]

which proves the claim by the direct comparison test for convergence of series.

Remark A.1. When Weyl law applies, we have that \( \lambda_\ell = \Theta(\ell^{2/n}) \), which allows us to replace the above condition by \( \sum \ell^{(n-1)/n} \alpha(\lambda_\ell) < \infty \). For the diffusion kernel/distance choice of \( \alpha(\lambda) = e^{-t\lambda} \) the series always converges independently of the manifold dimension. For biharmonic choice of \( \alpha(\lambda) = 1/\lambda^2 \), the sufficient condition is the convergence of \( \sum \ell^{(n-1)/\lambda} \sim \sum \ell^{(n-1)/(\ell^2/n)^2} = \sum \ell^{(n-5)/n} \), where we applied Weyl’s asymptotic again. As a result, the biharmonic choice of \( \alpha \) is guaranteed to provide a well-defined \( ISW_2 \) for 1 and 2-dimensional manifolds. Notice, however, that the Hörmander’s bound used in the proof of the above proposition can be rather lax in some of the settings that are practically relevant, such as the product spaces of lines and circles (where all of the eigenfunctions are bounded by a constant as can be seen from Table 4), and, thus, convergence for the biharmonic choice holds more widely.
Proposition 4.3. If \( D \) is a Hilbertian probability distance such that ISD is well-defined, then

(i) ISD is Hilbertian, and

(ii) ISD satisfies the following metric properties: non-negativity, symmetry, the triangle inequality, and \( ISD(\mu, \mu) = 0 \).

Proof. By Hilbertian property of \( D \), there exists a Hilbert space \( H^0 \) and a map \( \eta^0 : P(\mathbb{R}) \to H^0 \) such that \( D(\rho_1, \rho_2) = \|\eta^0(\rho_1) - \eta^0(\rho_2)\|_{H^0} \) for all \( \rho_1, \rho_2 \in P(\mathbb{R}) \). Plugging this into the definition of ISD we have \( ISD(\mu, \nu) = \|\eta(\mu) - \eta(\nu)\|_{H} \), where \( H = \oplus_{\ell} H^0 \) and the \( \ell \)-th component of \( \eta(\mu) \) is \( \sqrt{\alpha(\lambda_{\ell})} \eta_\ell(\phi_\ell^2 \mu) \in H^0 \). The second part of Proposition A.2 directly follows from the Hilbert property.

Proposition A.1. When \( \mu = \delta_x(\cdot), \nu = \delta_y(\cdot) \) for two points \( x, y \in \mathcal{X} \), we have \( ISW_2(\mu, \nu) = d(x, y) \), where \( d(\cdot, \cdot) \) is the spectral distance corresponding to the choice of \( \alpha(\cdot) \).

Proof. We have \( \phi_\ell \delta_x(x) = \delta_{\phi_\ell(x)}(\cdot) \) and similarly for \( y \). Now \( \mathcal{W}_2^2(\phi_\ell \sharp \mu, \phi_\ell \sharp \nu) = \mathcal{W}_2^2(\delta_{\phi_\ell(x)}, \delta_{\phi_\ell(y)}) = (\phi_\ell(x) - \phi_\ell(y))^2 \). This last equality follows from the fact that the 2-Wasserstein on real line between delta measures is equal to the distance between the two points. Then scaling and adding up gives exactly the kernel distance \( d(x, y) \) between the two points.

Proposition 4.4. Let \( D(\rho_1, \rho_2) = |E_{x\sim \rho_1}[x] - E_{y\sim \rho_2}[y]| \) for \( \rho_1, \rho_2 \in P(\mathbb{R}) \), then the corresponding intrinsic sliced distance is equivalent to the MMD with the spectral kernel \( k(\cdot, \cdot) \).

Proof. We can rewrite the definition as follows:

\[
ISD^2(\mu, \nu) = \sum_{\ell} \alpha(\lambda_\ell) (E_{x\sim \phi_\ell \sharp \mu}[x] - E_{y\sim \phi_\ell \sharp \nu}[y])^2 = \sum_{\ell} \alpha(\lambda_\ell) (|E_{x\sim \mu}[\phi_\ell(x)] - E_{y\sim \nu}[\phi_\ell(y)]|)^2
\]

\[
= \sum_{\ell} \alpha(\lambda_\ell) (E_{x, x' \sim \mu}[\phi_\ell(x)\phi_\ell(x')] + E_{y, y' \sim \nu}[\phi_\ell(y)\phi_\ell(y')] - 2E_{x \sim \mu, y \sim \nu}[\phi_\ell(x)\phi_\ell(y)])
\]

\[
= \sum_{\ell} \alpha(\lambda_\ell) E_{x, x' \sim \mu}[k(x, x')] + E_{y, y' \sim \nu}[k(y, y')] - 2E_{x \sim \mu, y \sim \nu}[k(x, y)],
\]

where we used the spectral kernel \( k(x, y) = \sum_{\ell} \alpha(\lambda_\ell) \phi_\ell(x)\phi_\ell(y) \). The last expression coincides with the MMD based on kernel \( k(\cdot, \cdot) \); see Lemma 6 in (Gretton et al., 2012).

Proposition 4.5. \( MMD(\mu, \nu) \leq ISW_2(\mu, \nu) \) when the same \( \alpha(\cdot) \) is used in both constructions.

Proof. This follows directly from the fact that for \( \rho_1, \rho_2 \in P(\mathbb{R}) \) the inequality \( |E_{x\sim \rho_1}[x] - E_{y\sim \rho_2}[y]| \leq \mathcal{W}_1(\rho_1, \rho_2) \leq \mathcal{W}_2(\rho_1, \rho_2) \) holds. Here the first inequality follows from the centroid bound (Rubner et al., 2000), and the second inequality is the well-known ordering property of Wasserstein distances (Villani, 2003).

Theorem 4.6. If \( \alpha(\lambda) > 0 \) for all \( \lambda > 0 \), then \( ISW_2 \) is a metric on \( P(\mathcal{X}) \).

Proof. In the light of the Proposition A.2 it remains only to prove that \( ISW_2(\mu, \nu) = 0 \) implies \( \mu = \nu \). According to Proposition A.2, \( ISW_2(\mu, \nu) = 0 \) yields \( MMD(\mu, \nu) = 0 \). The assumption that \( \alpha(\lambda) > 0 \) for all \( \lambda > 0 \) implies that the spectral kernel \( k(\cdot, \cdot) \) corresponding to \( \alpha(\cdot) \) is universal (Micchelli et al., 2006). Universality implies the characteristic property (Gretton et al., 2012), which in turn means that \( MMD(\mu, \nu) = 0 \) is equivalent to \( \mu = \nu \), proving the claim.

Proposition 4.7. There exists a constant \( c \) depending only on \( \mathcal{X} \) such that for all \( \mu, \nu \in P(\mathcal{X}) \) the inequality \( ISW_2(\mu, \nu) \leq c \mathcal{W}_2^X(\mu, \nu) \sqrt{\sum_{\ell} \lambda_\ell^{(n+3)/2}} \alpha(\lambda_\ell) \) holds; here, \( n \) is the dimension of \( \mathcal{X} \).

Proof. We remind \( \mathcal{W}_2^X \) is the 2-Wasserstein distance defined directly on \( P(\mathcal{X}) \) using the geodesic distance as the ground metric. The Neumann eigenfunctions on compact manifolds satisfy the inequality \( ||\nabla \phi_\ell||_\infty \leq c_1 \lambda_\ell ||\phi_\ell||_\infty \), see (Hu et al., 2015).
Applying the bound used in the proof of convergence, \( \|\phi_\ell\|_\infty \leq c_2 \lambda_\ell^{(n-1)/4} \), we get that \( \phi_\ell \) is Lipschitz with respect to the geodesic distance on \( \mathcal{X} \) with the Lipschitz constant bounded by \( c\lambda_\ell^{(n-1)/4} = c\lambda_\ell^{(n+3)/4} \).

Consider the optimal coupling between \( \mu \) and \( \nu \) whose cost equals to \( W_2^\mathcal{X}(\mu, \nu) \). Note that this coupling straightforwardly provides a coupling between the pushforwards \( \phi_\ell \mu \) and \( \phi_\ell \nu \). Using the Lipschitz property of eigenfunctions, we see that the cost of the pushforward coupling is smaller than \( c\lambda_\ell^{(n+3)/4} W_2^\mathcal{X}(\mu, \nu) \). Since any such coupling provides an upper bound on \( W_2(\phi_\ell \mu, \phi_\ell \nu) \), we have \( W_2(\phi_\ell \mu, \phi_\ell \nu) \leq c\lambda_\ell^{(n+3)/4} W_2^\mathcal{X}(\mu, \nu) \). Plugging this into the formula for \( ISW_2 \) we get the claimed bound.

**Proposition 4.8.** Let \( \{\mu_i\}_{i=1}^N \) and \( \{\nu_i\}_{i=1}^N \) be two collections of probability measures on \( \mathcal{P}(\mathcal{X}) \), such that \( \forall i, W_2^\mathcal{X}(\mu_i, \nu_i) \leq \epsilon \), then \( T(\{\mu_i\}_{i=1}^N, \{\nu_i\}_{i=1}^N) \leq C\epsilon^2 \). Here \( C = c\sqrt{\sum \lambda_\ell^{(n+3)/2}/\alpha(\lambda_i)} \) from previous proposition and is assumed to be finite.

**Proof.** We have

\[
T(\{\mu_i\}_{i=1}^N, \{\nu_i\}_{i=1}^N) = \left\| \frac{1}{N} \sum_{i=1}^N \eta(\mu_i) - \frac{1}{N} \sum_{i=1}^N \eta(\nu_i) \right\|_H^2 = \left\| \frac{1}{N} \sum_{i=1}^N (\eta(\mu_i) - \eta(\nu_i)) \right\|_H^2 \\
\leq \frac{1}{N} \sum_{i=1}^N \|\eta(\mu_i) - \eta(\nu_i)\|_H^2 = \frac{1}{N} \sum_{i=1}^N ISW_2^\mathcal{X}(\mu_i, \nu_i) \leq \frac{1}{N} \sum_{i=1}^N (C\lambda_\ell^{n/2}(\mu_i, \nu_i))^2 \\
\leq \frac{1}{N} (C\epsilon)^2 = C\epsilon^2. 
\]

**A.3. Computational Details for Section 4.2**

The case of finite intervals is the building block for the general case, so let us first consider the case of \( \mathcal{X} = [0, T] \). We represent a histogram over this interval by a discrete measure of the form \( \mu = \sum a \delta_{x_a} \) with the histogram bin centers \( x_a \in [0, T] \) and weights \( a \) satisfying \( \sum a = 1 \), where \( a = 1, 2, ..., A \). Note that it is not required for the histograms in the collections to be supported at the same bin locations. For a given histogram, let \( \{x_{(a)}, w_{(a)}\}_{a=1}^A \) be the locations sorted from smallest to largest and their corresponding weights; since the bin locations are unique there will not be any ties. The quantile function is computed via \( F^{-1}_\mu(s) := \min\{x_{(a)} : \sum_{b \leq a} w_{(b)} > s\} \). The approximate map \( \eta_{ID}^0 \) now can be computed using the \( s_k \)-th quantile value \( F^{-1}_\mu(s_k) \) for each value of \( s_k \), \( k = 1, ..., D' \).

For a general domain \( \mathcal{X} \), the histogram representation is the same as above: \( \sum a \delta_{x_a} \) with the histogram bin centers \( x_a \in \mathcal{X} \) and weights \( a \) satisfying \( \sum a = 1 \), where \( a = 1, 2, ..., A \). The pushforward \( \phi_\ell \mu \) gives a histogram on the real line defined by \( \sum \delta_{\phi(\phi(x_a))} \). Note that while \( x_a \) are distinct, their images under \( \phi_\ell \) do not have to be distinct, so one re-aggregates the weights to obtain \( \sum_{a \in S} w'_a \delta_{\phi(\phi(x_a))} \), where \( S \) is a subset of \( 1, 2, ..., A \) and \( w'_a \) are the new weights. It is now straightforward to compute the quantile function as before and build the approximate map \( \eta_{ID}^0 \). Doing so for the different values of \( \ell \) and concatenating the resulting vectors gives \( \eta_{ID} \).

In practice, these computations can be carried out on a variety of domains—analytic manifolds, manifolds discretized as point clouds or meshes, and graphs. In most cases the spectral decomposition of the Laplace-Beltrami operator or graph Laplacian has to be computed numerically (Coifman & Lafon, 2006; Reuter et al., 2009). For applications that involve simple manifolds, the eigenvalues and eigenfunctions can be computed analytically. For completeness we list them in Table 4. Note that we benefit from the fact that the eigen-decomposition for product spaces can be derived from the eigen-decompositions of the components.

The choice of the function \( \alpha(\cdot) \) determining the contributions of each spectral band is problem specific. When working on manifolds of low dimension, the choice of \( \alpha(\cdot) \) that corresponds to the biharmonic distance is convenient. While the diffusion distance provides a general choice that works on manifolds of any dimension, the biharmonic distance does not have any parameters to tune and was shown to provide an excellent alternative to the geodesic distance in low-dimensional settings (Lipman et al., 2010). When in doubt, inspecting the behavior of the distance on the underlying domain will allow assessing whether the distance is appropriate for the given problem. The importance of relying on a well-behaved spectral distance was highlighted in Proposition A.1.
Proof. Using the Hilbertianity of \( ISD \) (Proposition A.2), we have

\[
ISD^2(\mu_i, \mu_j) = \|\eta(\mu_i) - \eta(\mu_j)\|_{\mathcal{H}}^2 = \|\eta(\mu_i)\|_{\mathcal{H}}^2 + \|\eta(\mu_j)\|_{\mathcal{H}}^2 - 2\langle \eta(\mu_i), \eta(\mu_j) \rangle_{\mathcal{H}}.
\]

Consequently

\[
\sum_{i,j:i\neq j} ISD^2(\mu_i, \mu_j) = 2(N_1 - 1) \sum_{i=1}^{N_1} \|\eta(\mu_i)\|_{\mathcal{H}}^2 - 2 \sum_{i,j:i\neq j} \langle \eta(\mu_i), \eta(\mu_j) \rangle_{\mathcal{H}}.
\]
Similarly,  
\[
\sum_{i,j:i\neq j} ISD^2(\nu_i, \nu_j) = 2(N_2 - 1) \sum_{i=1}^{N_2} ||\eta(\nu_i)||_H^2 - 2 \sum_{i,j:i\neq j} \langle \eta(\nu_i), \eta(\nu_j) \rangle_H,
\]
\[
\sum_{i,j} ISD^2(\mu_i, \nu_j) = N_2 \sum_{i=1}^{N_1} ||\eta(\mu_i)||_H^2 + N_1 \sum_{j=1}^{N_2} ||\eta(\nu_j)||_H^2 - 2 \sum_{i,j:i\neq j} \langle \eta(\mu_i), \eta(\nu_j) \rangle_H.
\]

Putting these back into Eq. (6) after simplifying and cancelling out the norm-square terms we have
\[
\hat{T} = \frac{1}{N_1(N_1 - 1)} \sum_{i,j:i\neq j} \langle \eta(\mu_i), \eta(\mu_j) \rangle_H + \frac{1}{N_2(N_2 - 1)} \sum_{i,j:i\neq j} \langle \eta(\nu_i), \eta(\nu_j) \rangle_H - \frac{2}{N_1N_2} \sum_{i,j} \langle \eta(\mu_i), \eta(\nu_j) \rangle_H.
\]

At this point, we replace the maps \( \eta \) by their centered versions \( \tilde{\eta}(\mu) = \eta(\mu) - C_{\eta\#P}, \tilde{\eta}(\nu) = \eta(\nu) - C_{\eta\#Q} \); remember that the center of mass of \( \eta\#P \) is denoted by \( C_{\eta\#P} \). Accumulating the sample-level partial sums above the centering terms cancel out under \( H_0 : C_{\eta\#P} = C_{\eta\#Q} \), so that each \( \eta \) can be replaced by \( \tilde{\eta} \) in (9) above.

Denote \( x_i \equiv \tilde{\eta}(\mu_i), y_i \equiv \tilde{\eta}(\nu_i) \) as the Hilbert-embedded samples of \( X \sim \tilde{\eta}\#P, Y \sim \tilde{\eta}\#Q \), respectively. We remind now that \( R \) is a mixture of the centered pushforward measures: \( R = \rho_2(\tilde{\eta}\#P) + \rho_1(\tilde{\eta}\#Q) \). Let \( L_2(H, R) \) be the space of real-valued functions on \( H \) that are square integrable with respect to \( R \). Now we can define the following operator \( S : L_2(H, R) \rightarrow H \),
\[
(Sf)(x) := \int_H \langle x, x' \rangle_H f(x')dR(x').
\]

Following condition (ii), \( \langle \cdot, \cdot \rangle_H \) is square-integrable under \( R \). The above operator is thus Hilbert-Schmidt, hence compact (Reed & Simon, 1080, Theorem VI.23). Consequently, it permits an eigenfunction decomposition with respect to measure \( R, \langle x, x' \rangle_H = \sum_{m=1}^\infty \gamma_m \psi_m(x)\psi_m(x'), \) for \( x, x' \in H \). Note that here \( \psi_m : H \rightarrow \mathbb{R} \) and
\[
\int_H \langle x, x' \rangle_H \psi_m(x')dR(x') = \gamma_m \psi_m(x),
\]
\[
\int_H \psi_m(x)\psi_m(x)dR(x) = \delta_{mm}.
\]

Due to the centering of \( \eta \) we also have when \( \gamma_m \neq 0, \)
\[
\gamma_m \mathbb{E}_X[\psi_m(x)] = \int_H \mathbb{E}_X[\langle x, x' \rangle_H] \psi_m(x')dR(x') = 0 \Rightarrow \mathbb{E}_X[\psi_m(x)] = 0.
\]

Similarly, \( \mathbb{E}_Y[\psi_m(y)] = 0. \) The V-statistic from the overall sample can now be written as an infinite sum (Serfling, 2009, Section 5.5):
\[
\| \hat{C}_{\eta\#P} - \hat{C}_{\eta\#Q} \|_H^2 = \sum_{m=1}^\infty \gamma_m \left( \frac{1}{N_1} \sum_{i=1}^{N_1} \psi_m(x_i) - \frac{1}{N_2} \sum_{i=1}^{N_2} \psi_m(y_i) \right)^2 := \sum_{m=1}^\infty \gamma_m a_m^2.
\]

Our goal is to show that (a) \( a_m \sim N(0, (N_1\rho_1 N_2\rho_2)^{-1}) \), for \( \forall m \), and (b) \( a_m \) and \( a_n \) are independent when \( m \neq n \).

First note that
\[
\mathbb{E}(a_m) = \mathbb{E} \left( \frac{1}{N_1} \sum_{i=1}^{N_1} \psi_m(x_i) - \frac{1}{N_2} \sum_{i=1}^{N_2} \psi_m(y_i) \right) = 0.
\]
In addition we have,

\[
\text{Cov}(a_m, a_n) = \mathbb{E}(a_m a_n) - \mathbb{E}(a_m) \mathbb{E}(a_n) \\
= \mathbb{E}(a_m a_n) \\
= \mathbb{E}\left( \frac{1}{N_1} \sum_{i=1}^{N_1} \psi_m(x_i) - \frac{1}{N_2} \sum_{i=1}^{N_2} \psi_m(y_i) \right) \left( \frac{1}{N_1} \sum_{i=1}^{N_1} \psi_n(x_i) - \frac{1}{N_2} \sum_{i=1}^{N_2} \psi_n(y_i) \right) \\
= \mathbb{E}_X \left( \frac{1}{N_1} \sum_{i=1}^{N_1} \psi_m(x_i) \psi_n(x_i) \right) + \mathbb{E}_Y \left( \frac{1}{N_2} \sum_{i=1}^{N_2} \psi_m(y_i) \psi_n(y_i) \right) \\
= \frac{1}{\rho_1 N} \mathbb{E}_X \left( \frac{1}{N_1} \sum_{i=1}^{N_1} \psi_m(x_i) \psi_n(x_i) \right) + \frac{1}{\rho_2 N} \mathbb{E}_Y \left( \frac{1}{N_2} \sum_{i=1}^{N_2} \psi_m(y_i) \psi_n(y_i) \right) \\
= \frac{1}{\rho_1 \rho_2} \int_{\mathcal{H}} \psi_m(x) \psi_n(x) d(\tilde{\eta} \# P)(x) + \frac{1}{\rho_2} \int_{\mathcal{H}} \psi_m(y) \psi_n(y) d(\tilde{\eta} \# Q)(y) \\
= \frac{1}{\rho_1 \rho_2} \int_{\mathcal{H}} \psi_m(z) \psi_n(z) dR(z) \\
= \frac{1}{\rho_1 \rho_2} \delta_{mn}.
\]

An application of CLT follows that (a) holds. This together with vanishing covariance proves (b). Consequently, we can apply the CLT for degenerate V-statistics (Serfling, 2009, Section 5.5.2) to obtain the limiting distribution, with \( A_m \sim N(0, 1) \),

\[
N \| \tilde{C}_{\eta \# P} - \tilde{C}_{\eta \# Q} \|_{\mathcal{H}}^2 \sim \sum_{m=1}^{\infty} \frac{\gamma_m}{\rho_1 \rho_2} A_m^2.
\]

Let us now look at the difference between this V-statistic and our U-statistic, i.e. \( \hat{T} \) in (9). We see that

\[
\| \tilde{C}_{\eta \# P} - \tilde{C}_{\eta \# Q} \|_{\mathcal{H}}^2 - \hat{T} = \frac{1}{N_1} \sum_{i,j} \langle x_i, x_j \rangle_{\mathcal{H}} + \frac{1}{N_2} \sum_{i,j} \langle y_i, y_j \rangle_{\mathcal{H}} - \frac{2}{N_1 N_2} \sum_{i,j} \langle x_i, y_j \rangle_{\mathcal{H}} \\
- \frac{1}{N_1(N_1 - 1)} \sum_{i,j;i \neq j} \langle x_i, x_j \rangle_{\mathcal{H}} + \frac{1}{N_2(N_2 - 1)} \sum_{i,j;i \neq j} \langle y_i, y_j \rangle_{\mathcal{H}} + \frac{2}{N_1 N_2} \sum_{i,j} \langle x_i, y_j \rangle_{\mathcal{H}} \\
- \frac{1}{N_1(N_1 - 1)} - \frac{1}{N_2} \sum_{i,j;i \neq j} \langle x_i, x_j \rangle_{\mathcal{H}} + \left[ \frac{1}{N_2(N_2 - 1)} - \frac{1}{N_2^2} \right] \sum_{i,j;i \neq j} \langle y_i, y_j \rangle_{\mathcal{H}} \\
+ \left( \frac{1}{N_1} \sum_{i=1}^{N_1} \| x_i \|_{\mathcal{H}}^2 + \frac{1}{N_2} \sum_{i=1}^{N_2} \| y_i \|_{\mathcal{H}}^2 \right) \\
= -K^x - K^y + B.
\]

We claim that \( K^x = O_p(N_1^{-2}) \), \( K^y = O_p(N_2^{-2}) \), and \( NB \rightarrow P \sum_{m=1}^{\infty} \gamma_m (\rho_1 \rho_2)^{-1} \). As a result,

\[
N \left[ \| \tilde{C}_{\eta \# P} - \tilde{C}_{\eta \# Q} \|_{\mathcal{H}}^2 - \hat{T} \right] = -N O_p(N_1^{-2}) - N O_p(N_2^{-2}) + \sum_{m=1}^{\infty} \frac{\gamma_m}{\rho_1 \rho_2} + o_p(1) \\
= \sum_{m=1}^{\infty} \frac{\gamma_m}{\rho_1 \rho_2} + o_p(1),
\]

so that \( N \hat{T} \sim \sum_{m=1}^{\infty} \gamma_m (\rho_1 \rho_2)^{-1} (A_m^2 - 1) \), and we conclude the proof by reassigning \( \gamma_m \leftarrow \gamma_m (\rho_1 \rho_2)^{-1} \) to obtain (7).
Proof of Claim. For the $K$-terms we have

\[
K^x = \left[\frac{1}{N_1(N_1 - 1)} - \frac{1}{N_1^2}\right] \sum_{i,j \neq j} \langle x_i, x_j \rangle \mathcal{H}
\]

\[
= \frac{1}{N_1(N_1 - 1)} \sum_{i,j \neq j} \langle x_i, x_j \rangle \mathcal{H}
\]

\[
= \sum_{m=1}^{\infty} \gamma_m \frac{1}{N_1(N_1 - 1)} \sum_{i,j \neq j} \psi_m(x_i)\psi_m(x_j)
\]

\[
= \sum_{m=1}^{\infty} \gamma_m K_m^x,
\]

where $K_m^x$ is defined as the inner sum. Since $E\mathcal{X} \psi_m(x) = 0$, we have $E\mathcal{X}(K_m^x) = \frac{1}{N_1}[E\mathcal{X} \psi_m(x)]^2 = 0$, and

\[
Var_{\mathcal{X}}(K_m^x) = E\mathcal{X}[(K_m^x)^2]
\]

\[
= \frac{1}{N_1^2} E\mathcal{X} \left[ \frac{1}{N_1^2(N_1 - 1)} \sum_{i \neq j \neq k} \psi_m(x_i)\psi_m(x_j)\psi_m(x_k) \right]
\]

\[
= \frac{1}{N_1^2} E\mathcal{X} \left[ \frac{1}{N_1^2(N_1 - 1)} \sum_{i \neq j} \psi_m^2(x_i)\psi_m^2(x_j) \right]
\]

\[
= \frac{1}{N_1^2} \gamma_m (E\mathcal{X} \psi_m^2(x))^2.
\]

The cross terms—terms involving $l \neq i$ or $k \neq j$—vanish due to the sample being iid and eigenfunctions having zero expectations. The expectation in the last line is finite by assumption (ii), so that $Var_{\mathcal{X}}(K_m^x) = O(N_1^{-2})$, giving $K_m^x = O_p(N_1^{-2})$. Note that the assumption (ii) moreover implies the convergence of the big-oh coefficients, leading to $K^x = \sum_{m=1}^{\infty} \gamma_m K_m^x = O_p(N_1^{-2})$. Similarly we get $K^y = O_p(N_2^{-2})$.

For the term $B$, we have

\[
B = \frac{1}{N_1^2} \sum_{i=1}^{N_1} ||x_i||_\mathcal{H}^2 + \frac{1}{N_2^2} \sum_{i=1}^{N_2} ||y_i||_\mathcal{H}^2 = \sum_{m=1}^{\infty} \gamma_m \left[ \frac{1}{N_1^2} \sum_{i=1}^{N_1} \psi_m^2(x_i) + \frac{1}{N_2^2} \sum_{i=1}^{N_2} \psi_m^2(y_i) \right] := \sum_{m=1}^{\infty} \gamma_m C_m.
\]

Taking expectation,

\[
E_{\mathcal{X,Y}}(C_m) = \frac{1}{\rho_1 N_1} \int_\mathcal{H} \psi_m^2(x) d(\tilde{\mu} \# P)(x) + \frac{1}{\rho_2 N_2} \int_\mathcal{H} \psi_m^2(y) d(\tilde{\nu} \# Q)(y)
\]

\[
= \frac{1}{N_1 \rho_1 \rho_2} \int_\mathcal{H} \psi_m^2(z) dR(z)
\]

\[
= \frac{1}{N_1 \rho_1 \rho_2}.
\]

Thus $E_{\mathcal{X,Y}}(NB) = \sum_m \gamma_m (\rho_1 \rho_2)^{-1}$. Finally,

\[
NB = \sum_{m=1}^{\infty} \gamma_m \left[ \frac{1}{\rho_1 N_1} \sum_{i=1}^{N_1} \psi_m^2(x_i) + \frac{1}{\rho_2 N_2} \sum_{i=1}^{N_2} \psi_m^2(y_i) \right] \rho_1 \sum_{m=1}^{\infty} \gamma_m \left[ \frac{1}{\rho_1} E_{\mathcal{X}} \psi_m^2(x) + \frac{1}{\rho_2} E_{\mathcal{Y}} \psi_m^2(y) \right] = E_{\mathcal{X,Y}}(NB)
\]

by the weak law of large numbers. This proves the claim for $B$.

Alternative Distribution. For the the limiting distribution under $H_1$, notice that the first two terms in (6) are the one-sample U-statistic calculated on the samples $\{\mu_i\}_{i=1}^{N_1}$ and $\{\nu_i\}_{i=1}^{N_2}$, respectively. Using the CLT for non-degenerate U-statistics
We obtain (8) by combining the above three results.

As we mention in the discussion after condition (i), for the heat kernel with tuning parameter \( t \) (Serfling, 2009, Section 5.5.1, Theorem A), we have

\[
\hat \eta(c) = \text{fast and the finite dimensional embeddings can be constructed with small approximation errors, provided the manifold eigenvalues are declining suitably}
\]

where \( \delta, k > 0 \) are constants depending only on \( \mathcal{X} \).

The following result now ensures that approximations of \( \hat T \) using the top few eigenfunctions and a finite number of CDF embeddings can be constructed with small approximation errors, provided the manifold eigenvalues are declining suitably fast and the finite dimensional \( \eta_l(\cdot) \) is suitably smooth.

**Proposition A.3.** Suppose that (i), (ii) and (iii) hold. Then we have \( \sqrt{N} (\hat T - \hat T_{L_N}) = o_p(1) \) and \( \sqrt{N} (\hat T_{L_N} - \hat T_{L_N,D_N}) = o_p(1) \) for the following choices of \( L_N, D_N \):

\[
L_N \geq \min_{L'} \left\{ L': \sum_{\ell = L' + 1}^{\infty} \alpha_l \lambda_{l}^{(3/2)} \leq \frac{1}{N^{1+\delta}} \right\}, \quad D_N \geq k c^2 N^{1+\delta} \sum_{\ell=1}^{L_N} \alpha_l \lambda_{l}^{(n-1)/2},
\]

where \( \delta, k > 0 \) are constants depending only on \( \mathcal{X} \).

As we mention in the discussion after condition (i), for the heat kernel with tuning parameter \( t \): \( \alpha(\lambda) = \exp(-t \lambda) \), the assumption (i) that \( \sum_{\ell=1}^{\infty} \alpha_l \lambda_{l}^{(n+3)/2} < \infty \) holds. The bound on \( D_N \) is a consequence of classical bounds on Riemann sum approximation errors in terms of \( \|\eta'\|_{\infty} \). Absolute continuity of \( \mu \sim P, \nu \sim Q \) ensures the existence of \( (F_{\phi_{1\ell}})^{-1}(s), (F_{\phi_{2\ell}})^{-1}(s) \) (where prime denotes the derivative) for Lebesgue-almost every \( s \in [0,1] \) (Gavish et al., 2019, Lemma 2.3).

**Proof.** Notice that given \( L_N \), summands in the expression \( \hat T - \hat T_{L_N} \) are the tail sums \( \sum_{\ell=L_N+1}^{\infty} \alpha_l \lambda_{l}^{(n+3)/2} \) starting at the \( L_N + 1 \)th term. Using a similar approach as the proof of Proposition A.2, this is bounded above by a scalar multiple of the geodesic distance, specifically \( c \mathcal{W}_{2}^{X}(\cdot, \cdot) \sqrt{\sum_{\ell=L_N+1}^{\infty} \alpha_l \lambda_{l}^{(n+3)/2}} \). By assumption \( \sum_{\ell=1}^{\infty} \alpha_l \lambda_{l}^{(n+3)/2} < \infty \), so that given \( \epsilon > 0 \) we can always choose a starting point to make the tail sum < \( \epsilon \). The choice of \( L_N \) follows by taking \( \epsilon = N^{-(1+\delta)} \).

To obtain the choice of \( D_N \), we first use a similar approach to the proof of Proposition A.2 to simplify \( \hat T_{L,D'} \) for any \( L, D' \):

\[
\hat T_{L,D'} = \sum_{\ell=1}^{L} \left[ \frac{1}{N_1(N_1-1)} \sum_{i,j:i\neq j} \eta_{D'}^{(\phi_{1\ell} \mu_i)} \eta_{D'}^{(\phi_{1\ell} \mu_j)} + \frac{1}{N_2(N_2-1)} \sum_{i,j:i\neq j} \eta_{D'}^{(\phi_{2\ell} \nu_i)} \eta_{D'}^{(\phi_{2\ell} \nu_j)} \right] - \frac{2}{N_1 N_2} \sum_{i,j} \eta_{D'}^{(\phi_{1\ell} \mu_i)} \eta_{D'}^{(\phi_{2\ell} \nu_j)}. \tag{12}
\]

Recall that the inverse CDF transformation induced by \( \eta_{0}(\phi_{1\ell} \mu) = F_{\phi_{1\ell} \mu}^{-1} \) maps \([0,1]\) to a bounded interval that is the range of \( \phi_{1\ell} \), and \( \|\phi_{1\ell}\|_{\infty} \leq c \lambda_{\ell}^{(n-1)/4} \) using Hörmander’s bound on the supremum norm of the eigenfunctions. Using classical results on Riemann sum approximation errors (Bakhvalov, 2015; Brown & Steinberger, 2020) we thus have for any \( \ell \):

\[
\|\alpha(\eta_{0}(\phi_{1\ell} \mu), \eta_{1}(\phi_{1\ell} \nu))_{\mathcal{H}} - \eta_{D'}(\phi_{1\ell} \mu) \eta_{D'}(\phi_{2\ell} \nu)\|_{\infty} \leq \frac{k}{D'} \| (F_{\phi_{1\ell} \mu}^{-1} F_{\phi_{2\ell} \nu}^{-1}) \|_{\infty} \leq \frac{2k^2}{D'} \alpha_l \lambda_{\ell}^{(n-1)/2}.
\]
Given $L = L_N$, we simply choose $D' = D_N$ large enough to make the right hand side above smaller than $N^{-(1+\delta)}$. While it is possible to make the upper bound tighter using recent results (such as (Brown & Steinerberger, 2020)), the above coarser bound suffices for our purpose.

We now state a version of Theorem 5.1, with specifications for Theorem A.5.

**Theorem A.4.** Assume conditions (i)-(iii) hold. Define $N = N_1 + N_2$, and suppose that as $N_1, N_2 \to \infty$, we have $N_1/N \to \rho_1, N_2/N \to \rho_2 = 1 - \rho_1$, for some fixed $0 < \rho_1 < 1$. With $L \geq L_N, D' \geq D_N$ chosen per Proposition A.3, under $H_0 : C_{\eta#P} = C_{\eta#Q}$ we have

$$N\hat{T}_{L,D'} \sim \sum_{m=1}^{\infty} \gamma_m(A_m^2 - 1),$$

where $A_m, \gamma_m$ are defined as in Proposition A.2. Further, under $H_1 : C_{\eta#P} \neq C_{\eta#Q}$ we have $\sqrt{N}\left(\hat{T}_{L,D'} - T\right) \sim N(0, \sigma_1^2)$.

**Proof.** This a combination of Propositions A.2 and A.3, and Slutsky’s theorem.

We conclude with a proof of a specified version of Theorem 5.2, which gives power guarantee of the test based on $\hat{T}_{L,D'}$ for contiguous alternatives.

**Theorem A.5.** Assume conditions (i)-(iii) hold, and let $L, D'$ be chosen as in Theorem A.4. Then for the sequence of contiguous alternatives $H_{1N}$ such that $N\|\delta_N\|_H^2 \to \infty$, the test based on $\hat{T}_{L,D'}$ is consistent for any $\alpha \in (0, 1)$, that is as $N \to \infty$ the asymptotic power approaches 1.

**Proof.** It is enough to prove consistency using $\hat{T}$, as the difference between $\hat{T}$ and $\hat{T}_{L,D'}$ is negligible by choice of $L, D'$. To do so we utilize proof techniques similar to Theorem 13 in Gretton et al. (2012). Define $c_N := N^{1/2}\|\delta_N\|_H$, and expand the simplified centered version of the test statistic in (9) but under $H_1$ so that the centering terms do not cancel out:

$$\hat{T}_c = \frac{1}{N_1(N_1 - 1)} \sum_{i,j:i\neq j} \langle \eta(\mu_i) - C_{\eta#P}, \eta(\mu_j) - C_{\eta#P} \rangle_H$$

$$+ \frac{1}{N_2(N_2 - 1)} \sum_{i,j:i\neq j} \langle \eta(\nu_i) - C_{\eta#Q}, \eta(\nu_j) - C_{\eta#Q} \rangle_H$$

$$- \frac{2}{N_1N_2} \sum_{i,j} \langle \eta(\mu_i) - C_{\eta#P}, \eta(\nu_j) - C_{\eta#Q} \rangle_H.$$ (13)

The centered pushforwards have the same Hilbert centroids, thus as $N \to \infty$ by Proposition A.2,

$$N\hat{T}_c \sim \sum_{m=1}^{\infty} \gamma_m(A_m^2 - 1) := S.$$ (14)

Subtracting $\hat{T}_c$ from $\hat{T}$ and its expansion in Eq. (6) on the left and right hand respectively, then simplifying we have

$$N(\hat{T} - \hat{T}_c) = N \left[ -\frac{1}{N_1} \sum_{i=1}^{N_1} \langle \delta_N, \eta(\mu_i) - C_{\eta#P} \rangle_H + \frac{1}{N_2} \sum_{i=1}^{N_2} \langle \delta_N, \eta(\nu_i) - C_{\eta#Q} \rangle_H + \frac{\langle \delta_N, \delta_N \rangle_H}{2} \right]$$

$$= N \left[ -\frac{\|\delta_N\|_H}{N_1} \sum_{i=1}^{N_1} \langle \delta_N, \eta(\mu_i) - C_{\eta#P} \rangle_H$$

$$- \frac{\|\delta_N\|_H}{N_2} \sum_{i=1}^{N_2} \langle \delta_N, \eta(\nu_i) - C_{\eta#Q} \rangle_H + \frac{\|\delta_N\|_H^2}{2} \right].$$ (14)
Given \( N \) the inner products \( \langle \delta_N / ||\delta_N||_H, \eta(\mu_i) - C_{\eta\#P} \rangle_H \) are i.i.d. random variables with mean 0, so by CLT then using \( ||\delta_N||_H = c_N N^{-1/2} \) we get

\[
\frac{1}{\sqrt{N_2}} \sum_{i=1}^{N_2} \left\langle \frac{\delta_N}{||\delta_N||_H}, \eta(\mu_i) - C_{\eta\#P} \right\rangle_H \sim U \quad \Rightarrow \quad \frac{N ||\delta_N||_H}{N_1} \sum_{i=1}^{N_2} \left\langle \frac{\delta_N}{||\delta_N||_H}, \eta(\nu_i) - C_{\eta\#Q} \right\rangle_H \sim \frac{c_N}{\sqrt{p_1}} U, 
\]

where \( U \) is the zero mean Gaussian random variable that is the limiting distribution of the above inner product sum. Similarly we have

\[
\frac{N ||\delta_N||_H}{N_2} \sum_{i=1}^{N_2} \left\langle \frac{\delta_N}{||\delta_N||_H}, \eta(\nu_i) - C_{\eta\#Q} \right\rangle_H \sim \frac{c_N}{\sqrt{p_2}} V, 
\]

where \( V \) is also Gaussian, zero mean, and independent of \( U \). Putting everything together in the right hand side of (14), and using \( ||\delta_N||_H = c_N N^{-1/2} \), given the threshold \( t_\alpha \) for a level-\( \alpha \) test

\[
P_{H_N} \left( N T > t_\alpha \right) \rightarrow P \left[ S + c_N \left( \frac{U}{\sqrt{p_1}} - \frac{V}{\sqrt{p_2}} \right) + \frac{c_N^2}{2} > t_\alpha \right].
\]

By assumption \( c_N^2 \rightarrow \infty \), so the asymptotic power approaches 1 as \( N \rightarrow \infty \).

\[\square\]

A.5. Proofs and Notes for Section 5.2

To guarantee size control when using the the harmonic mean \( p \)-value we establish a version of Theorem 1 from Liu & Xie (2020). Assume that a test statistic \( Z \in \mathbb{R}^D \) has null distribution with zero mean and every pair of coordinates of \( Z \) follows bivariate Gaussian distribution. Compute the coordinate-wise two-sided \( p \)-values \( p_k = 2(1 - \Phi(|Z_k|)) \) where \( \Phi \) is the standard Gaussian CDF.

**Theorem 5.4.** Let \( p_k, k = 1, \ldots, D \) be the null \( p \)-values as above and \( p^H \) computed via harmonic mean approach, then

\[
\lim_{\alpha \rightarrow 0} \frac{\text{Prob}\{ p^H \leq \alpha \}}{\alpha} = 1.
\]

**Proof.** The proof of Theorem 1 from (Liu & Xie, 2020) hinges on Lemma 3 in their supplemental material. We show that Lemma 3 holds for the harmonic mean combination method. Note that the multiplication by \( \pi \) present in Lemma 3 cancels out when inverse cotangent with a multiplier of \( 1/\pi \) is applied later on; so it is not relevant to the flow of the proof.

To this end, consider the functions \( p(x) = 2(1 - \Phi(|x|)) \) and \( h(x) = 1/p(x) \). We need to prove the following three statements:

1. For any \( |x| > \Phi^{-1}(3/4) \),

\[
\frac{\cos[p(x)\pi]}{p(x)} \leq h(x) \leq \frac{1}{p(x)}
\]

2. For any constant \( 0 < |a| < 1 \), we have

\[
\lim_{x \rightarrow +\infty} \frac{h(x)}{x^2 h(ax)} > c_a > 0,
\]

where \( c_a \) is some constant only dependent on \( a \).

3. Suppose that \( X_0 \) has standard normal distribution, then we have

\[
P\{ h(X_0) \geq t \} = \frac{1}{t} + O(1/t^3).
\]

Statement (1) is trivial, as \( h(x) = 1/p(x) \) by definition and the cosine function is upper bounded by one. Statement (2) holds by the same argument as in the supplement of (Liu & Xie, 2020). Statement (3) follows from the fact that when \( X_0 \) is standard normal, then \( p(x) \) is a null \( p \)-value, and so

\[
P\{ h(X_0) \geq t \} = P\{ p(X_0) \leq 1/t \} = \frac{1}{t}.
\]
We also compare the performance of our tests on data from a number of domains with several existing methods, and settings of the embedding parameters \(L, D\). For evaluation, we use empirical power at different degrees of departure from the null hypothesis, calculated by averaging the proportion of rejections at level \(\alpha = 0.05\) over 1000 independent datasets with samples divided into two groups of sizes \(n_1 = 60, n_2 = 40\). To ensure the tests are well-calibrated, we also calculate nominal sizes assuming the two sample groups are drawn from the same random meta-distribution. We calculate eigenvalues and eigenfunctions using analytical expressions provided in Table 4. We fix \(\lambda(t) = e^{-t}\) (i.e. heat kernel with \(t = 1\)) for all experiments, in order to avoid unfair advantage from tuning this parameter when comparing to baselines. Also, when using the \(p\)-value combination test each \(t\)-test scales the mean difference by standard deviation, so the weight functions cancel out anyway. In general, when \(t\) is small, the \(T\) statistic is more democratic between low and high frequency eigenfunctions, allowing to capture finer details of the underlying domain (assuming large enough \(L\)). When \(t\) is large, \(T\) focuses on low frequency eigenfunctions so more global differences dominate the computation of \(T\).

**Finite intervals** To obtain our base measures \(\mu_i, \nu_i\), we generate bin probabilities as (shifted and normalized) values of the function \(f(t_j) = \mu(t_j) + \alpha(t_j)\) at \(m = 30\) fixed design points \(t_j = j/(m+1), j = \{1, 2, \ldots, m\}\), and

\[
\begin{align*}
\mu(t_j) &= 1.2 + 2.3 \cos(2\pi t_j) + 4.2 \sin(2\pi t_j), \\
\alpha(t_j) &= \epsilon_0 + \sqrt{2}\epsilon_1 \cos(2\pi t_j) + \sqrt{3}\epsilon_2 \sin(2\pi t_j),
\end{align*}
\]

where \(\epsilon_0, \epsilon_1, \epsilon_2 \sim N(0, 1)\) clipped between \([-3, 3]\). Group 1 and 2 samples are obtained as \(\mu_i(\cdot) \equiv f(\cdot)\) and \(\nu_i(\cdot) \equiv f(\cdot) + \delta\) respectively, where \(\delta \in [0, 4]\) is a constant. To make the sample functions non-negative, we shift all functions by \(M = 3(1 + \sqrt{2} + \sqrt{3})\). Finally, as the \(m\)-length vector of bin counts for a sample, we generate a random vector from the Multinomial distribution with 1000 trials, \(m\) outcomes and the outcome probabilities proportional to the shifted functional observations corresponding to that sample.

We use embedding dimensions \(L = 3, D' = 10\) to compare our method against 11 functional ANOVA tests—for brevity we report results for 3 of them which use different methodological approaches (see Appendix for complete results). All methods maintain nominal size for \(\delta = 0\) (Figure 3 a). While the combination test (ISD comb) based on our proposal outperformed all the other tests across all values of \(\delta\), the bootstrap test that uses the overall \(T\) statistic (ISD T boot) performs better than Fmaxb but worse than others. Table 5 shows the outputs for the other 8 competing methods from the R package \(fdANOVA\) for the finite intervals synthetic data setting.\(^1\)

We also compare the \(p\)-value combination test based on an unsliced 24-dimensional inverse CDF embedding with sliced \(ISW_2\)-based tests (Figure 3 b). We use multiple pairs of \((L, D')\) values, all of them giving overall embeddings of dimension \(D = LD' = 24\). The performance of an \(ISW_2\)-based test that uses slicing over only the first eigenfunction is almost as good as the unsliced version. With more eigenfunctions, the powers first improve considerably, then become similar to the unsliced version again.

\(^1\)See https://www.rdocumentation.org/packages/fdANOVA/versions/0.1.2/topics/fanova.tests for full names of all methods.
Intrinsic Sliced Wasserstein Distances on Manifolds and Graphs

Figure 4. Performance on synthetic finite interval and manifold data. Finite interval: (a) comparison with existing methods—a test based on basis function representation (FP) (Gorecki & Smaga, 2015), a sum-type $\ell_2$ norm-based test (L2b) (Zhang, 2013), and a max-type test (Zhang et al., 2019) that uses the maximum of coordinate-wise $F$ statistic (Fmaxb); (b) unsliced vs. different settings of $(L, D')$. Manifold data: (c) circular data, comparing with Fréchet ANOVA (Dubey & Müller, 2019), and the DISCO nonparametric test (Rizzo & Székely, 2010); (d) harmonic combination tests on cylindrical data for $L = 4$. Dotted lines indicates nominal size of all tests ($\alpha = 0.05$).

| $\delta$ | CH | CS | L2N | L2b | FN | FB | Fb | GPF |
|---------|----|----|-----|-----|----|----|----|-----|
| 0       | 0.031 | 0.03 | 0.033 | 0.024 | 0.031 | 0.028 | 0.033 | 0.026 |
| 0.1     | 0.025 | 0.024 | 0.03 | 0.044 | 0.027 | 0.03 | 0.041 | 0.021 |
| 0.2     | 0.026 | 0.029 | 0.037 | 0.06 | 0.033 | 0.034 | 0.058 | 0.025 |
| 0.3     | 0.036 | 0.041 | 0.044 | 0.067 | 0.041 | 0.04 | 0.067 | 0.033 |
| 0.4     | 0.034 | 0.035 | 0.036 | 0.057 | 0.034 | 0.035 | 0.056 | 0.032 |
| 0.5     | 0.051 | 0.052 | 0.058 | 0.091 | 0.056 | 0.057 | 0.088 | 0.044 |
| 0.6     | 0.056 | 0.066 | 0.066 | 0.089 | 0.061 | 0.066 | 0.088 | 0.051 |
| 0.7     | 0.07 | 0.083 | 0.083 | 0.121 | 0.084 | 0.081 | 0.119 | 0.064 |
| 0.8     | 0.085 | 0.097 | 0.095 | 0.151 | 0.093 | 0.094 | 0.144 | 0.081 |
| 0.9     | 0.118 | 0.142 | 0.14 | 0.2 | 0.144 | 0.137 | 0.194 | 0.118 |
| 1       | 0.158 | 0.182 | 0.176 | 0.232 | 0.183 | 0.173 | 0.228 | 0.154 |
| 1.1     | 0.215 | 0.247 | 0.246 | 0.303 | 0.251 | 0.242 | 0.301 | 0.212 |
| 1.2     | 0.27 | 0.31 | 0.303 | 0.375 | 0.311 | 0.3 | 0.368 | 0.27 |
| 1.3     | 0.328 | 0.363 | 0.357 | 0.438 | 0.37 | 0.353 | 0.43 | 0.324 |
| 1.4     | 0.395 | 0.432 | 0.432 | 0.504 | 0.436 | 0.423 | 0.499 | 0.394 |
| 1.5     | 0.488 | 0.52 | 0.514 | 0.592 | 0.521 | 0.511 | 0.586 | 0.483 |
| 1.6     | 0.534 | 0.595 | 0.576 | 0.652 | 0.593 | 0.566 | 0.647 | 0.544 |
| 1.7     | 0.628 | 0.677 | 0.669 | 0.723 | 0.678 | 0.661 | 0.719 | 0.631 |
| 1.8     | 0.704 | 0.737 | 0.727 | 0.789 | 0.748 | 0.725 | 0.785 | 0.707 |
| 1.9     | 0.785 | 0.823 | 0.812 | 0.869 | 0.827 | 0.806 | 0.867 | 0.793 |
| 2       | 0.83 | 0.849 | 0.844 | 0.88 | 0.85 | 0.841 | 0.875 | 0.832 |
| 2.1     | 0.865 | 0.888 | 0.881 | 0.916 | 0.887 | 0.878 | 0.915 | 0.872 |
| 2.2     | 0.903 | 0.922 | 0.916 | 0.946 | 0.928 | 0.912 | 0.946 | 0.907 |
| 2.3     | 0.938 | 0.95 | 0.944 | 0.964 | 0.951 | 0.944 | 0.963 | 0.944 |
| 2.4     | 0.958 | 0.973 | 0.967 | 0.977 | 0.972 | 0.966 | 0.976 | 0.964 |
| 2.5     | 0.974 | 0.98 | 0.976 | 0.985 | 0.981 | 0.975 | 0.985 | 0.974 |
| 2.6     | 0.977 | 0.981 | 0.979 | 0.987 | 0.981 | 0.978 | 0.986 | 0.977 |
| 2.7     | 0.989 | 0.996 | 0.992 | 0.997 | 0.996 | 0.992 | 0.997 | 0.991 |
| 2.8     | 0.997 | 0.998 | 0.997 | 0.998 | 0.998 | 0.997 | 0.998 | 0.996 |
| 2.9     | 0.996 | 0.997 | 0.996 | 0.999 | 0.997 | 0.996 | 0.999 | 0.997 |
| 3       | 0.998 | 1 | 0.999 | 1 | 1 | 0.999 | 1 | 0.999 |

Table 5. Outputs for other methods in the functional curves synthetic data setting.

Manifold domains We consider data from distributions on circles and cylinders. For circular data, we take von Mises distributions with randomly chosen parameters as our samples. For an angle $x$ (measured in radians), the von Mises probability density function is given by $f(x|\mu, \kappa) = \exp[\kappa \cos(x - \mu)]/(2\pi I_0(\kappa))^{-1}$, where $I_0(\kappa)$ is the modified Bessel
function of order 0. We fix $\kappa = 2$, and use $\mu \equiv \mu_i \sim N(0, 0.1^2)$, $\mu \equiv \nu_i \sim N(\delta, 0.1^2)$ for samples from group 1 and 2 respectively—with $\delta \in [0, 15] \times \pi/180$ (i.e. 0 to 15 degrees converted to radians). As each observation vector, we take 100 random draws from each sample-specific distribution. For our embeddings, we use $L = 10$, $D' = 20$, and so our final embedding dimension is $10 \times 20 \times 2 = 400$. Since the competing methods cannot handle circular geometry directly, to implement them we cut the circle into an interval. Figure 3 (c) shows that all methods maintain nominal size, but both our tests maintain considerably higher power than existing methods for all $\delta$.

We generate cylindrical data in the form of samples of a bivariate random vector $(\Theta, X)$, using the cylindrical density function proposed by (Mardia & Sutton, 1978):

$$f(\theta, x) = \frac{e^{\kappa \cos(\theta - \mu)}}{2\pi I_0(\kappa)} \frac{1}{\sqrt{2\pi \sigma_c}} \frac{e^{-\frac{(x - \mu_c)^2}{2\sigma_c^2}}}{},$$

clipping values of the $X$-coordinate between the bounded interval $[0, 2\pi]$. This distribution has the parameters $\mu \in [-\pi, \pi], \mu_0 \in \mathbb{R}, \kappa \geq 0, \rho_1 \in [0, 1], \rho_2 \in [0, 1], \sigma > 0$, where $\mu, \kappa$ denote parameters for the (circular) marginal along the $\Theta$-coordinate. and given $\Theta = \theta, X$ is sampled from $N(\mu_c, \sigma_c^2)$, with

$$\mu_c = \mu + \sqrt{\kappa \sigma_c} \left\{ \rho_1 (\cos \theta - \cos \mu) + \rho_2 (\sin \theta - \sin \mu) \right\},$$

$$\sigma_c = \sigma^2 (1 - \rho^2), \rho = (\rho_1^2 + \rho_2^2)^{1/2}.$$

In our experiments, we fix $\rho_1 = \rho_2 = 0.5, \sigma = 1, \kappa = 2$ across both populations. As random samples of distributions, we draw $\mu, \mu_0 \sim \text{Unif}(0, 1)$ and $\mu, \mu_0 \sim \text{Unif}(\delta, \delta + 1)$ for samples of group 1 and 2 respectively, with $n_1 = 60, n_2 = 40$. We repeat the above for $\delta \in [0, 30]$ degrees converted to radians, and obtain bivariate histograms corresponding to each sample distribution from 500 random draws from that distribution. To evaluate the effects of choosing $L, D'$ we calculate our embeddings for $L \in \{2, 3, 4, 5\}, D' \in \{6, 8, 12, 24, 48\}$. The choice of $L$ has small effect on performance, so we report results for $L = 4$ in Figure 3 (d). Higher values of $D'$ result in some increase in power.

**Discussion** Our ISW$_2$-based method is able to exploit the non-euclidean nature of the problems and and their generality beyond mean comparison more effectively than competing methods, which are based on mean comparison on functional data/densities (frechet ANOVA, all functional ANOVA methods), and/or L2 distance-based comparisons (all functional ANOVA methods, DISCO). Regarding the optimal choice of embedding dimensions, while proving theorem 5.1 we show that (Proposition 10 therein) choosing both $L$ and $D$ above certain thresholds ensures close approximation to the population test statistic. For the combination test, adding more dimensions to the embedding can have a two-fold effect: a) probing more dimensions can help with finding differences, but b) every dimension adds another test and so potentially leads to loss of power. Thus, for the combination test, there must be an optimal data dependent choice of the embedding dimension, which can potentially be found via split testing procedures. We leave this to future work.

**Computational Complexity** Assume an underlying graph $G = (V, E)$, and we use $L$ slices, $D'$ quantiles to calculate ISW$_2$. Each distribution consists of $V$ atoms (distribution lives on graph vertices). Then, computation of our embeddings includes the following steps:

1. The computation of eigenvectors/values of symmetric sparse matrices is a well-studied problem with stable and efficient algorithms available, e.g. ARPACK providing an implementation of the Arnoldi method in MATLAB, Python, or R. These methods incur linear time complexity in the size of graph: $O(L(|V| + |E|))$. This computation is done only once per domain, and the overhead is negligible ($<< 1$ second) for our graph experiments.

2. Hilbert embedding computations require computing $L$ pushforwards and $D'$ quantiles, which need sorting. This complexity is $O(L(|V| \log |V| + |D'|))$.

3. Testing using p-value combination requires computing t-test p-values for $LD'$ dimensions, with overall complexity of $O(LD'N)$ where $N$ is total sample size.

In contrast, Sobolev Transport (ST) requires computing the Sliced ST Distance, based on computing shortest paths from randomly selected nodes in a graph. This has complexity $O(k(|V| \log |V| + |E|))$ (Le et al., 2022, pg. 4) Implementing permutation tests means repeating the above computation $P$ times, with $P = 1000$ being a typical choice.
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Figure 5. Activity histograms for three individuals from NHANES dataset. There are 100 bins in the intensity and 96 in the time dimension; we show hour of day on the time axis. The time dimension is periodic where 00:00 is identified with 24:00, giving rise to a cylindrical histogram domain.

Figure 6. Three eigenfunctions for the NHANES histogram domain normalized by the maximum absolute value. Note that the eigenfunctions are periodic in the time direction (i.e. match when glued over the side cut) but not in the intensity direction, reflecting the cylindrical geometry of the underlying domain.

B.2. NHANES data on physical activity monitoring

As our first real data application, we analyze the Physical Activity Monitor (PAM) data from the 2005-2006 National Health and Nutrition Examination Survey (NHANES)\textsuperscript{2}. This contains physical activity pattern readings for a large number of people collected over 1 week period on a per-minute granularity. After basic pre-processing steps to ensure no missing entries, as well as data reliability and well-calibrated activity monitors, we use data from 6839 individuals. The data for each individual corresponds to device intensity value from the PAM for $24 \times 60 = 1440$ minutes throughout the day, for 7 days.

For each individual we can capture their activity patterns into a cylindrical histogram with time and intensity dimensions. For each observation, its time during the day is discretized into 15-minute intervals giving 96 bins for the time dimension; its intensity value (capped at 1,000) is discretized into a 100 equidistant bins. Since the time dimension is periodic, we obtain a histogram over the cylinder $S^1(T_1) \times [0, T_2)$, with $T_1 = 96$, $T_2 = 100$. Normalized counts can thus be considered as person-specific probability distributions; several examples are shown in Figure 5. Note that flattening the domain by cutting the cylinder will arbitrarily split activity patterns (see especially Figure 5, Female 37) and will lead to inefficiencies due to horizontal variability.

We apply the proposed methodology to check if the activity patterns vary across different groups of individuals obtained as follows. We first split the overall dataset based on the individual’s age using the following inclusive ranges: 6–15, 16–25, ..., 76–85; this covers all the ages in the dataset. From each split we sample 100 males and 100 females to avoid gender imbalance driving the results. Thus, we end up with 8 age groups with 200 individuals per group. Our goal is to compare

\textsuperscript{2}https://wwwn.cdc.gov/Nchs/Nhanes/2005-2006/PAXRAW_D.htm
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Table 6. Comparing the activity intensity of different age groups based on the NHANES dataset. Below diagonal: $p$-values corresponding to the actual data comparisons. Above diagonal: null $p$-values obtained by combining and randomly splitting the two involved groups. The entries in boldface correspond to the rejected hypotheses with the BH procedure at the FDR level of 0.1.

| Age Groups | 6–15 | 16–25 | 26–35 | 36–45 | 46–55 | 56–65 | 66–75 | 76–85 |
|------------|------|-------|-------|-------|-------|-------|-------|-------|
| 6–15       | 0.979| 0.31  | 0.383 | 0.297 | 0.905 | 0.921 | 0.326 |
| 16–25      | 0.998| 0.963 | 0.443 | 0.872 | 0.442 | 0.529 |
| 26–35      | 0.987| 0.818 | 0.93  | 0.731 | 0.992 |
| 36–45      | 0.945| 0.984 | 0.974 | 0.327 |
| 46–55      | 0.377| 0.832 | 0.618 | 0.844 |
| 56–65      | 0.977| 0.987 | 0.818 | 0.93  |
| 66–75      | 0.977| 0.987 | 0.818 | 0.93  |
| 76–85      | 0.977| 0.987 | 0.818 | 0.93  |

To perform our analysis we compute the eigenvalues and eigenfunctions as per the 4th row of Table 4 using $\ell_1 = 1, 2, 3$ and $\ell_2 = 1, 2, 3$, giving a total of $L = 2 \times 3 \times 3 = 18$ eigenfunctions; three of the resulting eigenfunctions are shown in Figure 6. We consider a $D' = 5$ dimensional embedding for the inverse CDF transformation, hence the final embedding dimension after the slicing construction is $D = LDL' = 18 \times 5 = 90$.

We summarize the results in Table 6, below the diagonal. The $p$-values are obtained via the harmonic mean combination approach. We run the Benjamini-Hochberg (Benjamini & Hochberg, 1995) procedure on the resulting $p$-values at the false discovery rate of 0.1, and the rejected hypotheses are indicated by the $p$-values in bold. Our method detects statistically significant differences between all pairs of groups, except 46–55 versus 36–45 and 56-65 groups. As a control experiment, we provide our method with null cases and display the $p$-values in Table 6, above the diagonal. The null cases are obtained by combining the individuals from the two comparison groups and splitting it arbitrarily (i.e. mixing the two age groups). As expected, the $p$-values of the control comparisons do not concentrate near zero.

Curiously, our method can be used “off-label” to conduct functional data analyses over different dimensions of the NHANES dataset. For example, one can concentrate on a single day of activity intensity data which gives a curve over the 24-hour circle. Since activity intensity is a non-negative number, these curves can be normalized so as to obtain probability distributions. Now we can use our methodology to detect pair-wise differences across groups. While this has the benefit of accounting for underlying geometry of data, it loses the absolute magnitude information due to the normalization. Clearly the appropriateness of such an analysis would depend on the goal of the exercise and the particular research question attached to that goal; our proposal provides a framework that is flexible enough to handle data of different modalities.

### B.3. Chicago Crime

We demonstrate the use of our methodology on histograms over graphs. In this experiment, we use the Chicago Crimes 2018 dataset\(^3\) which captures incidents of crime in the City of Chicago. We base our analysis on the type of crime, the beat (geographic area subdivision used by police, see Figure 7) where the incident took place, and the date of the incident.

To capture the spatial aspect of the data we build a graph with one vertex per beat; two vertices are connected by an edge if the corresponding beats share a geographic boundary. For each crime type and day, we capture the total count of that crime type for each beat; after normalizing this gives a daily probability distribution over the graph. Our goal is to compare the collection of distributions of, say, theft occurring on Tuesday to those of Thursday and Saturday. The Tuesday versus Thursday comparison is intended as a null case, as we do not expect to see any differences between them (Rustamov & Klosowski, 2020).

We build the un-normalized Laplacian of the beat adjacency graph, and compute its lowest frequency $L = 20$ eigenvalues and eigenvectors. The first three eigenvectors are plotted in Figure 7. The number of inverse CDF values used in the embedding is $D' = 5$, which gives rise to $D = 100$ dimensional embedding. The results of comparisons are shown in the last two columns of Table 7; the $p$-values are obtained via the harmonic mean combination approach. We run the Benjamini-Hochberg (Benjamini & Hochberg, 1995) procedure on the 20 resulting $p$-values at the false discovery rate of 0.1, and the rejected hypotheses are indicated by the $p$-values in bold. As expected, no differences were detected between

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\(^3\)data.cityofchicago.org
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Figure 7. First three eigenvectors of the Laplacian are shown for the beat adjacency graph, mapped back to the geographic locations. All of the eigenvectors are normalized by the maximum absolute value. The spatial smoothness of the eigenvectors—somewhat masked here due to the discrete colormap—is crucial to efficiently capturing horizontal variability of the data (i.e. distribution shifts over the graph). The boundaries of beats are shown based on the shape file from Chicago Data portal.

Table 7. Results on Chicago Crime 2018 dataset. The entries in bold correspond to the rejected hypotheses with the BH procedure at the FDR level of 0.1. The \( N \) column captures the number of days passing the filtering criteria, and the \( \text{count} \) column shows the average per-day crime count.

| Crime Type            | Tuesday | Thursday | Saturday | Tue vs Thu | Tue vs Sat | p-value | p-value |
|-----------------------|---------|----------|----------|------------|------------|---------|---------|
| Theft                 | 52      | 178.7    | 52       | 182.9      | 52         | 180.2   | 0.452   | 4.7e-06 |
| Deceptive Practice    | 51      | 55.8     | 52       | 54.9       | 52         | 44.4    | 0.255   | 4.2e-04 |
| Battery               | 52      | 125.8    | 52       | 123.0      | 52         | 154.9   | 0.374   | 0.001   |
| Robbery               | 50      | 25.2     | 50       | 25.1       | 52         | 28.1    | 0.130   | 0.002   |
| Narcotics             | 51      | 36.0     | 51       | 34.6       | 50         | 36.9    | 0.890   | 0.008   |
| Criminal Damage       | 52      | 70.0     | 52       | 73.7       | 52         | 83.0    | 0.901   | 0.03    |
| Other Offense         | 52      | 49.5     | 52       | 48.4       | 52         | 44.1    | 0.670   | 0.037   |
| Burglary              | 52      | 34.0     | 52       | 33.1       | 52         | 29.1    | 0.157   | 0.183   |
| Motor Vehicle Theft   | 52      | 27.9     | 52       | 26.2       | 51         | 28.1    | 0.923   | 0.365   |
| Assault               | 52      | 57.2     | 52       | 59.3       | 52         | 52.4    | 0.996   | 0.617   |

Tuesday and Thursday patterns. On the other hand, we see that there are statistically significant differences between Tuesday and Saturday patterns in the following categories of crime: theft, deceptive practice, battery, robbery, narcotics, and criminal damage.

B.4. Brain Connectomics

In this example, we consider two publicly available brain connectomics datasets (Aine et al., 2017; Arroyo Relión et al., 2019) distributed as a part of the R package graphclass\(^4\). Both are based on resting state functional magnetic resonance imaging (fMRI): COBRE has data on 54 schizophrenics and 70 controls, and UMich with 39 schizophrenics and 40 controls. The datasets capture the pairwise correlations between 264 regions of interest (ROI) of Power parcellation (Power et al., 2011) and can be considered as a 264 node graph (263 nodes for COBRE as ROI 75 is missing) with positive and negative edge weights.

We define three probability measures supported on the nodes of the graph. For each ROI we take the sum of absolute values of all its correlations with the remaining ROIs. Now we have a positive number assigned to each node capturing its overall connectivity to the rest of the graph and we normalize to obtain a measure; this construction will be referred to as “all correlations”. Note that each scanned subject gives rise to a separate “all correlations” probability measure on the same

\(^4\)http://github.com/jesusdaniel/graphclass

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| Dataset | All correlations | Positive correlations | Negative correlations |
|---------|------------------|-----------------------|-----------------------|
| COBRE   | 0.0084           | 0.00019               | 0.0019                |
| UMich   | 0.609            | 0.116                 | 0.022                 |

Table 8. Comparison results between the schizophrenic and control groups for brain connectomics datasets.

underlying node set. The “positive correlations” and “negative correlations” constructions are based on keeping respectively only positive or only negative correlations and aggregating as above.

We also need a fixed base graph for the computation of the Laplacian eigen-decomposition; this graph should capture the spatial connectivity of the ROIs which is relevant due to the smooth nature of the blood oxygenation level dependent (BOLD) signal that is used for computing the correlations. To this end, we obtain the coordinates for the centers of the 264 ROIs\(^5\) and build the base graph by connecting each ROI to its nearest 8 ROIs. We compute the lowest frequency \(L = 20\) eigenvalues and eigenvectors of the corresponding un-normalized Laplacian. The number of inverse CDF values used in the embedding is \(D' = 5\), which gives rise to \(D = 100\) dimensional embedding.

Table 8 shows the result of comparing the schizophrenic group to the control group for both of the datasets; the \(p\)-values are obtained via the harmonic mean combination approach. We can see that our approach detects statistically significant differences between the two groups in COBRE dataset in all of the three types of measures on graphs. In contrast, for UMich dataset, the difference is detected only in the negative correlations and loses significance when corrected for multiple testing. This is potentially caused by the higher inhomogeneity of the UMich dataset that was pooled across five different experiments spanning seven years (Arroyo Relión et al., 2019). An interesting aspect of our analysis is that due to normalization (to obtain probability measures) the total sum of connectivity is factored out by the proposed method. As a result, the detected differences are not related to the well-known change in the overall connectivities between the two groups, but rather to distributional changes in marginal connectivity strengths.

\(^{5}\)www.jonathanpower.net/2011-neuron-bigbrain.html