Quantum Computation in a Topological Data Analysis Pipeline

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Topology in data



Large Data Sets



Main goal of Topological Data Analysis (TDA) Find and quantify structure in noisy, complex data. 1 Persistent Homology and Wasserstein Distance



Section 1

Persistent Homology and Wasserstein Distance



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Double annulus example



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Wasserstein QUBO

Double annulus example



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Definition

A distance on a set M is a function $d: M \times M \rightarrow \mathbb{R}_{\geq 0}$ such that

• $d(x, y) \ge 0$ and d(x, y) = 0 iff x = y

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$$d(x,y) = d(y,x)$$

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$$d(x,y) + d(y,z) \ge d(x,z)$$



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Wasserstein distance for diagrams

Given diagrams X and Y, the distance between them is

$$d_p(X, Y) = \inf_{\varphi: X \to Y} \left(\sum_{x \in X} \|x - \varphi(x)\|^p \right)^{1/p}$$



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Example



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Computation of Wasserstein Distance





 $\omega(u,v) = \|u-v\|^p$

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The **cost** of a matching $\mathbf{x} \subseteq E$ is the sum of the weights,

$$C_p(\mathbf{x}) = \sum_{e \in \mathbf{x}} \omega(e) = \sum_{(u,v) \in \mathbf{x}} \|u - v\|^p.$$



Equivalence

Theorem (Edelsbrunner et al 2010) $d_p(X, Y) = \varepsilon$ iff $\varepsilon^p = \min\{C(\mathbf{x}) \mid \mathbf{x} \text{ is a maximal matching}\}.$



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Definition

A min-cost maximal matching (MCMM) is a maximal matching ${f y}$ for which

 $C(\mathbf{y}) = \min\{C(\mathbf{x}) \mid \mathbf{x} \text{ is a maximal matching}\}$



Section 2

A Qubo for Wasserstein distance

The variables

$$\begin{array}{c} \mathbf{x} \in (\mathbb{Z}_2)^{nm+n+m} \\ \text{Binary variables:} \quad \mathbf{x} = \{x_{u,v} \mid (u,v) \in E\} \\ & & \\ & \\ & & \\ & \\ \text{Sets of edges} \qquad & \mathbf{x} \subset E \end{array}$$



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Warning: Not just maximal matchings



The QUBO





Theorem (Berwald, Gottlieb, EM, 2018) Assume $B > B^* := \max_{(u,v) \in E(\widetilde{G})} \omega(u, v)$. Then **x** is a solution which minimizes Hif and only if $\mathbf{x} \subseteq E$ is a MCMM of \widetilde{G} . Theorem (Berwald, Gottlieb, EM, 2018) Assume $B > B^* := \max_{(u,v) \in E(\widetilde{G})} \omega(u, v)$. Then **x** is a solution which minimizes Hif and only if $\mathbf{x} \subseteq E$ is a MCMM of \widetilde{G} .

Proof sketch

- For every subset z, there is a matching y with H(y) < H(z).
- For every non-maximal matching y there is a maximal matching x with H(x) < H(y).
- Maximal matchings have $H(\mathbf{x}) = C(\mathbf{x})$

Experiment







Experiment







Results



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- Why do the tests fail for larger sized problems?
 - Long chains of physical qubits?
- Other places in the persistent homology pipeline that can be swapped out for QC
 - Flag/Clique complexes
 - Computation of full persistent homology (Betti numbers: Lloyd et al 2016, Siopsis 2018, Dridi Alghassi 2015)
 - Multiparameter persistence

Thank you!

Relevant papers

- J. Berwald, J. Gottlieb, EM. *Computing Wasserstein Distance for Persistence Diagrams on a Quantum Computer*. arXiv:1809.06433, 2018
- EM. A User's Guide to Topological Data Analysis. Journal of Learning Analytics, 2017







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