

A Fast Method of Computing Persistent Homology of Time Series Data

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Introduction

- **Persistent homology** is applied to **reconstructed attractors** from time series data.
 - For periodicity detection and classification of time series data.
- Topological structure of orbits of ordinary differential equations reflects the periodicity or chaos.
- Computing persistent homology of thousands of points requires several days and hundreds GiB of memory.
 - We want to reduce the computational cost by fitting **cubic Bézier curves** to time series data.

Background

- Persistent homology extracts the number and the widths of holes in a shape.
- A shape is given as a simplicial complex.
- **Vietoris-Rips complex** is constructed from a point cloud.

Persistent homology

- For a point cloud X , the Vietoris-Rips complex of X is defined as

$$VR_r(X) = \{x_{i_0}x_{i_1} \cdots x_{i_q} \mid q\text{-simplex} \mid \forall j, k, d(x_{i_j}, x_{i_k}) \leq r\}.$$

- The simplices in $VR_r(X)$ is sorted by the value of the filter function:

$$f(\sigma) = \begin{cases} 0, & (\dim \sigma = 0), \\ d(u, v), & (\dim \sigma = 1), \\ \max_{\tau \prec \sigma} f(\tau), & (\text{otherwise}). \end{cases}$$

The Number of Simplices in a filtration of Vietoris-Rips complex

- The threshold r is set to the infinity.
- The upper dimension of homology groups is 2, so we need simplices of dimension less than 3.
- The number of simplices is $\sum_{k=0}^2 \binom{n}{k}$ and its order is $O(n^3)$ where n is the number of points.

Persistence Diagrams and Bottleneck Distance

- The **persistence diagram** of a persistent homology $PH_q(X)$ is defined as:

$$Dgm_q(X) = \{(b(\alpha), d(\alpha)) \mid \alpha \in PH_q(X)\} \cup \{(x, x) \mid x \in \mathbb{R} \cup \{\infty\}\}.$$

- The **bottleneck distance** between diagrams X and Y is defined as:

$$W_\infty(X, Y) = \inf_{\eta: X \rightarrow Y} \sup_{\alpha \in X} \|\alpha - \eta(\alpha)\|_\infty,$$

where $\eta: X \rightarrow Y$ ranges over all bijections and the norm is defined as

$$\|\alpha\|_\infty = \|(b, d)\|_\infty = \max\{|b|, |d|\}.$$

Attractor reconstruction

- The observed time series is mapped into the **delay-coordinate space**.
- The delay-coordinate vector with dimension d and delay a is defined as

$$y(t) = (x(t), x(t-a), x(t-2a), \dots, x(t-(d-1)a))$$

for a time series data $\{x(t)\}_{t=1}^n$.

Acknowledgements

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Proposed Method

Fit cubic Bézier curves

- The series of pairs of points and sampling time:

$$\{(x_i, t_i) \mid x_i \in \mathbb{R}^d\}_{i=1}^l \quad (0 \leq t_i \leq 1).$$

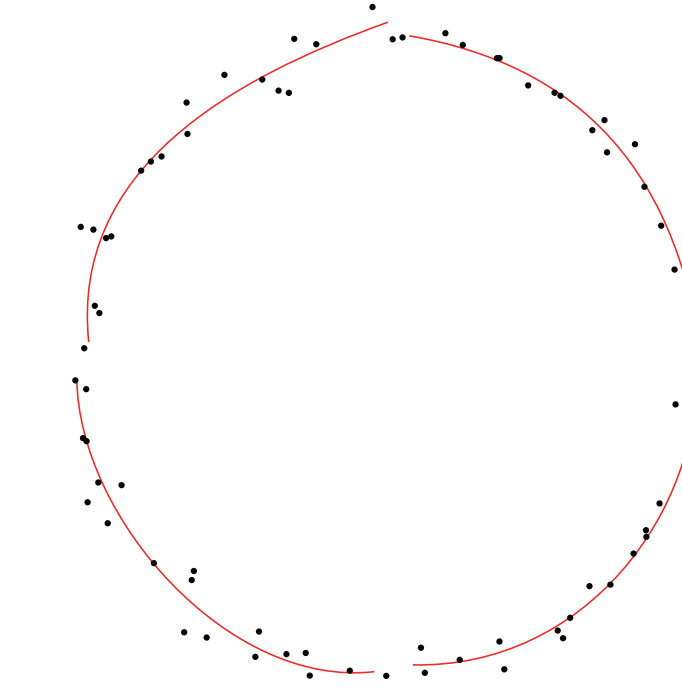
- The input time series is grouped into several groups.

- The cubic Bézier curve is parametrized as

$$p(t) = \sum_{i=0}^3 \binom{3}{i} (1-t)^{3-i} t^i p_i \quad (0 \leq t \leq 1).$$

- Fitting is achieved by minimizing the squared error function:

$$L(p_0, p_1, p_2, p_3) = \sum_{i=1}^l \|p(t_i) - x_i\|^2.$$



Divide into line segments

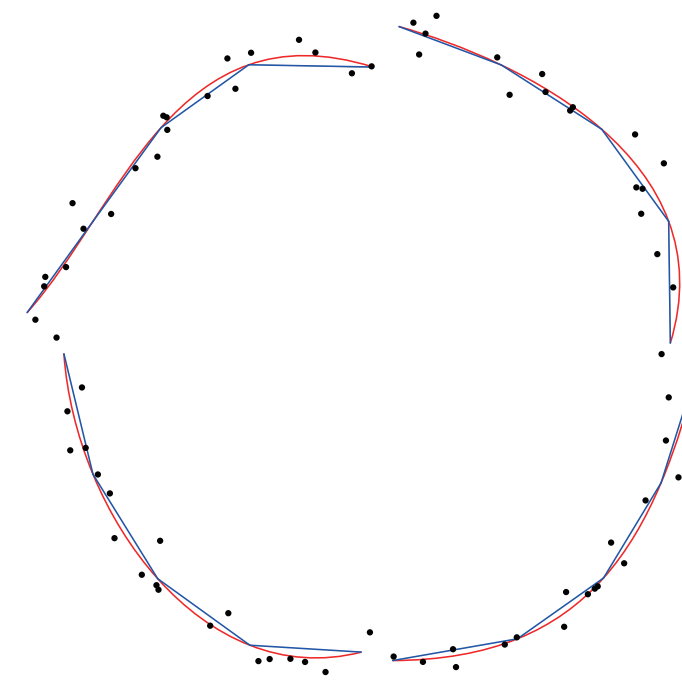
- The fitted cubic Bézier curve is divided into line segments.

- The unit interval $[0, 1]$ is uniformly divided into r intervals:

$$[t_0, t_1], \dots, [t_{r-1}, t_r] \quad \text{where } t_i = i/r \quad (i = 0, 1, \dots, r).$$

- Each interval is mapped into the cubic Bézier curve:

$$\overrightarrow{p(t_0)p(t_1)}, \overrightarrow{p(t_1)p(t_2)}, \dots, \overrightarrow{p(t_{r-1})p(t_r)}.$$



Distance between line segments

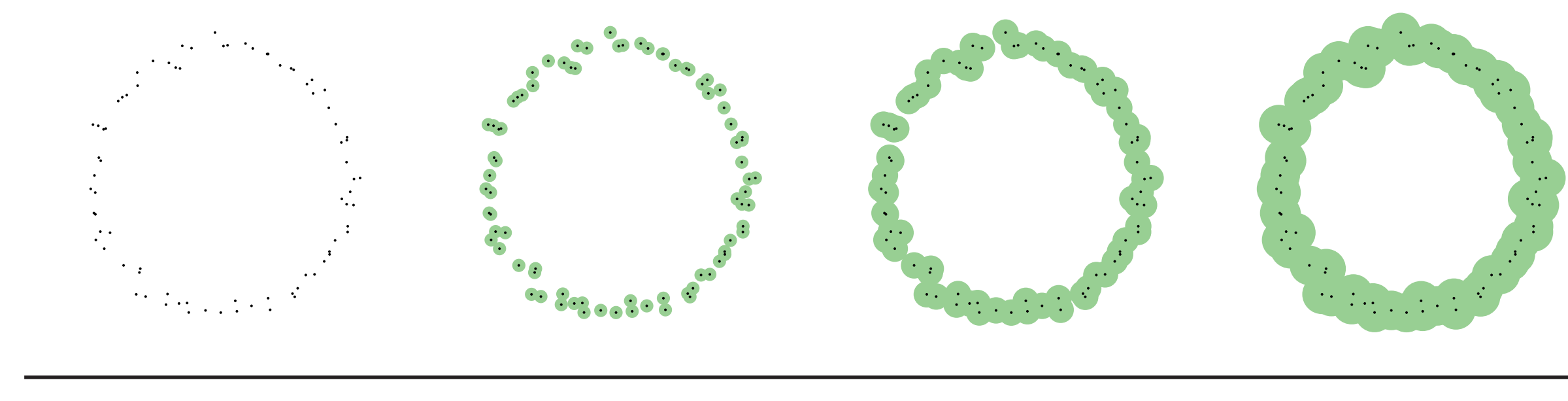
- Let $\overrightarrow{q_0q_1}$ and $\overrightarrow{r_0r_1}$ be line segments.

- The distance between them is defined as $d(\overrightarrow{q_0q_1}, \overrightarrow{r_0r_1}) = \min_{q \in \overrightarrow{q_0q_1}, r \in \overrightarrow{r_0r_1}} d(q, r)$.

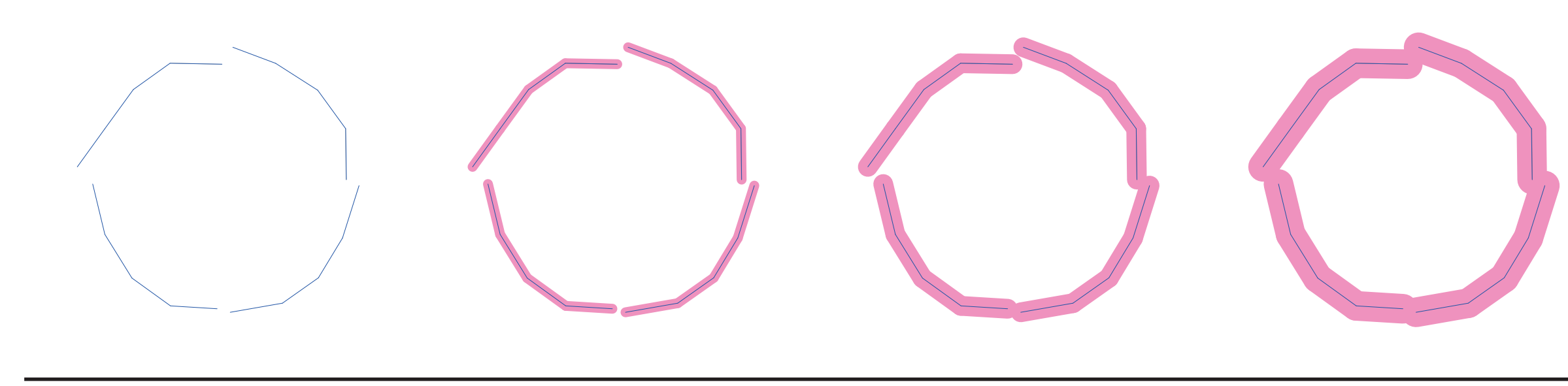
- The distance is calculated by minimizing the function $f(s, t) = d(q(s), r(t))^2 = \|q(s) - r(t)\|^2$.

Comparison between the ordinary and proposed Method

Ordinary Method



Proposed Method



Experiments

Irrational flow on 2-torus

- The irrational flow is defined as $\frac{du}{dt} = \alpha \pmod{1}$, $\frac{dv}{dt} = \beta \pmod{1}$, where $(u, v) \in [0, 1] \times [0, 1]$.

- We set $\alpha = 1$ and $\beta = \sqrt{2}$.

- The trajectory is mapped into 3-dimensional Euclidean space with the mapping below:

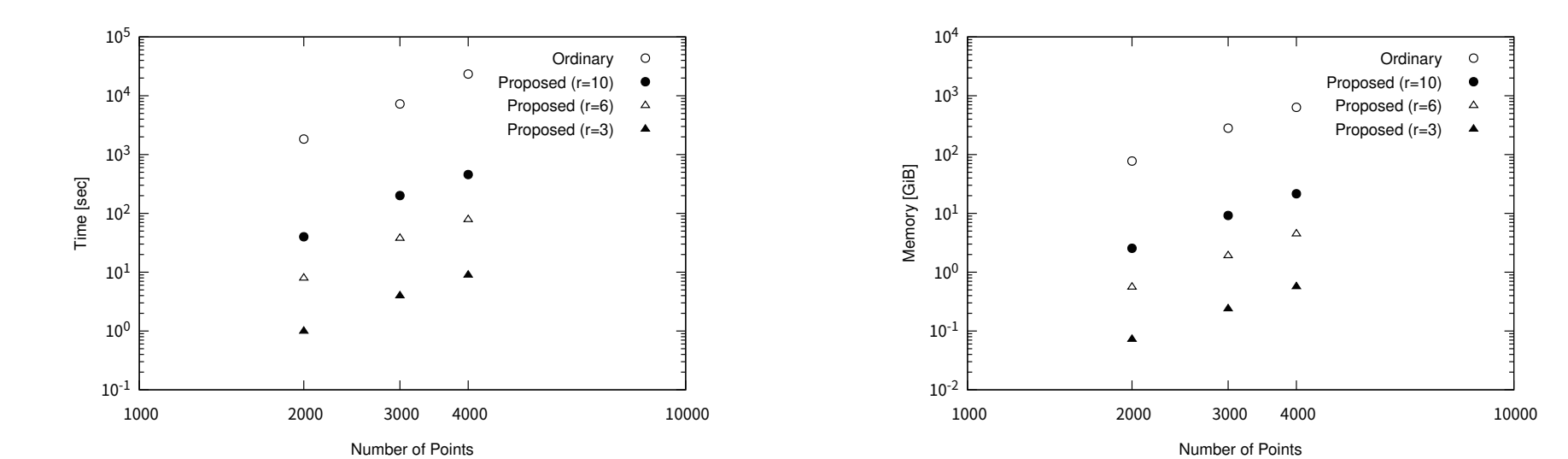
$$\begin{aligned} x_1 &= R \cos u + r \cos u \cos v, \\ x_2 &= R \sin u + r \sin u \cos v, \\ x_3 &= r \sin v, \end{aligned}$$

where $R = 2$ and $r = 1$.

- The trajectory was developed from $t = 0$ to $t = 50\pi$.

- The number of points was $n = 2000, 3000, 4000$ and the parameters was $l = 30$ and $r = 3, 6, 10$.

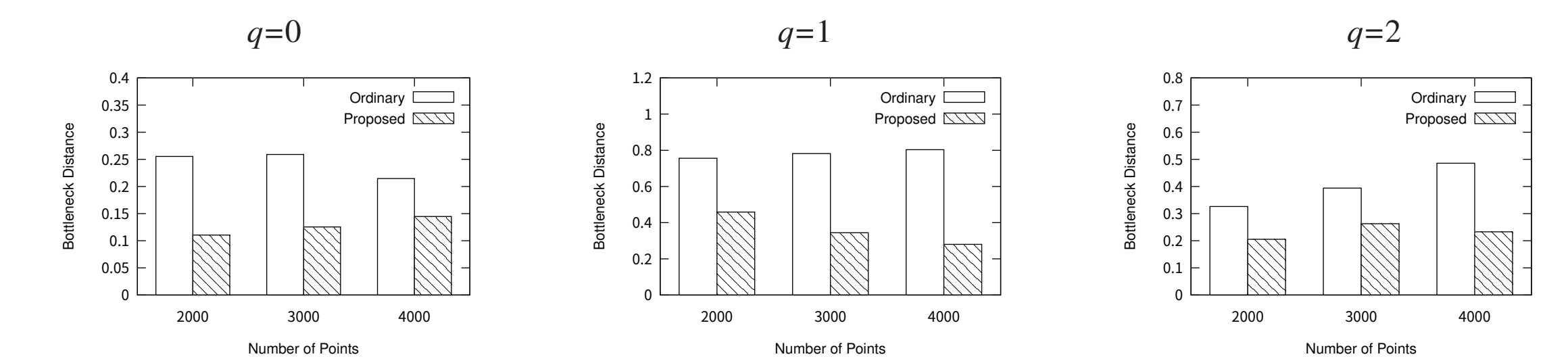
Time performance (left) and memory performance (right)



Comparison for noised input data

- Added 10% gaussian noise to the trajectory.

- Computed the bottleneck distance between the clean input and the noised input.



Japanese vowels /a/, /e/, /i/, /o/ and /u/

- The signals were mapped into the delay-coordinate space of $d = 10$ and $a = 10$.

- 1100 steps (almost 125 ms) of each embedded signal were extracted.

- The parameter was set to $l = 10$ and $r = 2$.

- The performance of the proposed method was compared to that of the **Witness complex**.

Vowel	Computational Time			Vowel	Computational Memory		
	Ordinary	Proposed	Witness		Ordinary	Proposed	Witness
/a/	290 sec	< 1 sec	< 1 sec	/a/	12.5 GiB	0.11 GiB	0.14 GiB
/e/	312 sec	< 1 sec	< 1 sec	/e/	11.5 GiB	0.09 GiB	0.09 GiB
/i/	268 sec	< 1 sec	< 1 sec	/i/	11.5 GiB	0.09 GiB	0.10 GiB
/o/	291 sec	< 1 sec	< 1 sec	/o/	11.5 GiB	0.09 GiB	0.09 GiB
/u/	277 sec	< 1 sec	< 1 sec	/u/	11.5 GiB	0.09 GiB	0.09 GiB

- The distances to the ordinary Vietoris-Rips complex was compared.

Vowel	Proposed Method			Vowel	Witness complex		
	0th	1st	2nd		0th	1st	2nd
/a/	0.064	0.644	0.131	/a/	0.126	0.645	0.177
/e/	0.029	0.032	0.047	/e/	0.026	0.217	0.038
/i/	0.026	0.027	0.014	/i/	0.018	0.292	0.034
/o/	0.021	0.068	0.011	/o/	0.022	0.279	0.069
/u/	0.016	0.032	0.020	/u/	0.015	0.332	0.029