

# SKELETON CLUSTERING AND REGRESSION.

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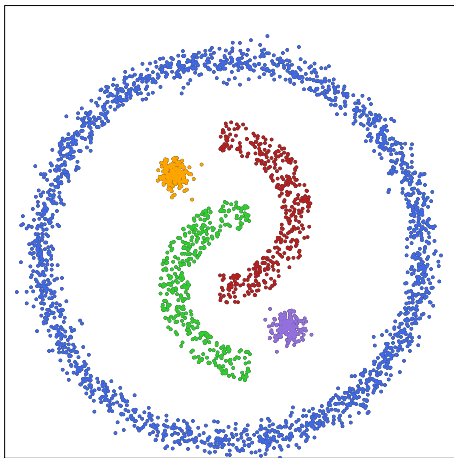
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Joint work with Jerry (Zeyu) Wei

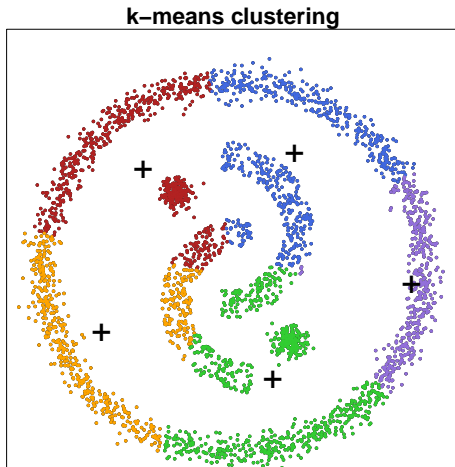


# A simple clustering problem in $d=1000$



Data:  $d = 1000$ ; only first 2 coordinates are shown here and the rest coordinates are Gaussian noises.

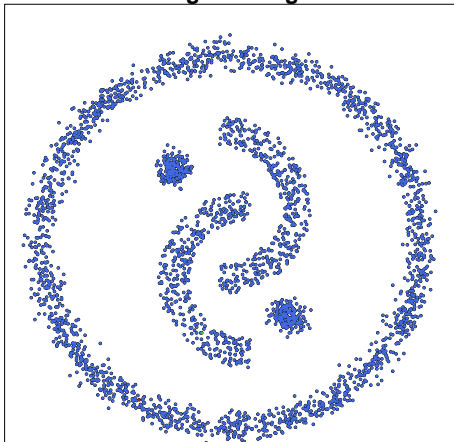
# A simple clustering problem in $d=1000$



Data:  $d = 1000$ ; only first 2 coordinates are shown here and the rest coordinates are Gaussian noises.

# A simple clustering problem in $d=1000$

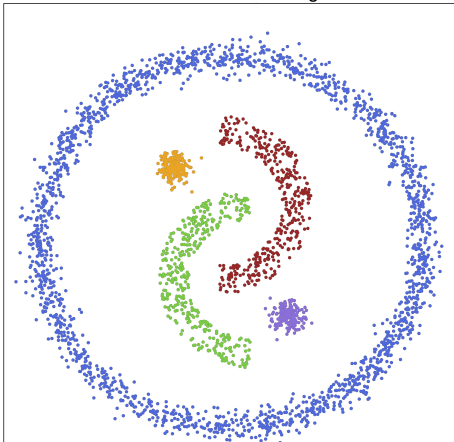
**Single Linkage**



Data:  $d = 1000$ ; only first 2 coordinates are shown here and the rest coordinates are Gaussian noises.

# A simple clustering problem in $d=1000$

Skeleton Clustering



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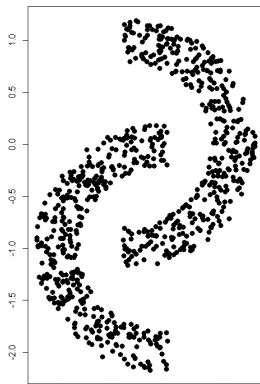
# Skeleton Clustering-1

Idea:

- We start with applying k-means clustering to the whole data with a large  $k$ .
- We then merge two clusters if they overlap a lot.

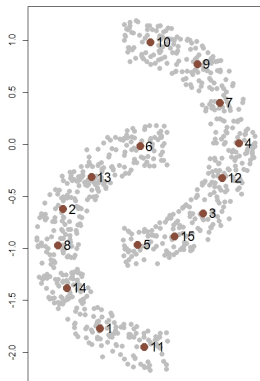
This procedure can be summarized as constructing a weighted graph called skeleton graph.

## Skeleton Clustering - 2



Original data.

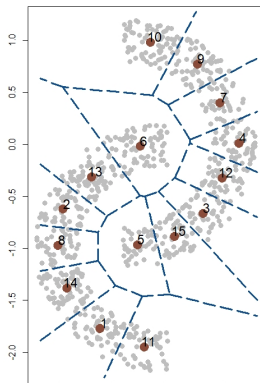
## Skeleton Clustering - 2



$k$ -means: generating knots (centers of  $k$ -means clusters);  $k = \sqrt{n}$ .

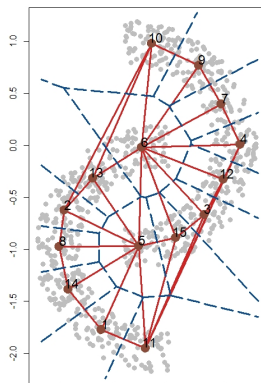


## Skeleton Clustering - 2



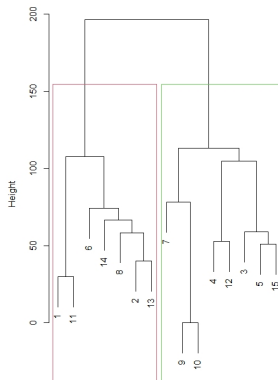
Voronoi cells of knots.

## Skeleton Clustering - 2



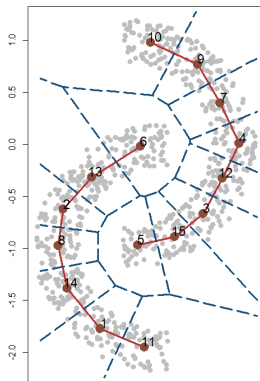
Delaunay triangulations—creating a graph. Assign a density-based weights on the edge to measure overlapping.

## Skeleton Clustering - 2



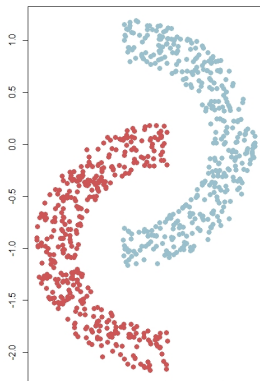
Convert a weighted graph into a dendrogram.

## Skeleton Clustering - 2



Cut the dendrogram to form the final clusters.

## Skeleton Clustering - 2



Final clustering result.

# Measuring the overlap between two knots

- Data:  $X_1, \dots, X_n \in \mathbb{R}^d$ .
- Centers of k-means:  $c_1, \dots, c_k$ .
- Partition of data:

$$\mathcal{X}_\ell = \{X_i : d(X_i, c_\ell) < d(X_i, c_j), \quad j \neq \ell\}$$

for knot  $c_\ell$ .

- Goal: we want to create a quantity to measure the overlap between  $c_j, c_\ell$ .

# Voronoi density

- Intuition:  $c_j$  and  $c_\ell$  have a high overlap if
  1. they are close, i.e.,  $\|c_j - c_\ell\|$  is small,
  2. there are many observations between the  $c_j$  and  $c_\ell$
- Define the 2-NN region of  $c_j$  and  $c_\ell$ :

$$A_{j\ell} = \{x : d(x, c_k) > \max\{d(x, c_j), d(x, c_\ell)\}, \forall k \neq j, \ell\}.$$

Namely, the 2-NN of knots at  $x$  is  $c_j, c_\ell$ .

- Let  $\mathbb{P}_n(A) = \frac{1}{n} \sum_{i=1}^n I(X_i \in A)$  be the empirical measure of the set  $A$ .
- We define the Voronoi density of  $c_j, c_\ell$  as

$$\widehat{S}_{j\ell}^{VD} = \frac{\widehat{P}_n(A_{j\ell})}{\|c_j - c_\ell\|}.$$

- Recall that

$$A_{j\ell} = \{x : d(x, c_k) > \max\{d(x, c_j), d(x, c_\ell)\}, \forall k\},$$
$$\widehat{S}_{j\ell}^{VD} = \frac{\widehat{P}_n(A_{j\ell})}{\|c_j - c_\ell\|}.$$

- Clearly,  $\widehat{S}_{j\ell}^{VD} = 0$  if  $c_j$  and  $c_\ell$  do not share a boundary.
- A population version of  $\widehat{S}_{j\ell}^{VD}$  is

$$S_{j\ell}^{VD} = \frac{P(A_{j\ell})}{\|c_j - c_\ell\|},$$

where  $P(A_{j\ell}) = P(X_i \in A_{j\ell})$ . The convergence is fast assuming that knots are fixed.



## Alternatives: Face density

- The Voroni density is not the only way to measure the overlap.
- We also have some other good alternatives although the Voronoi density works the best in practice.
- Face density:

$$\widehat{S}_{j\ell}^{FD} = \widehat{\rho}_{j\ell} \left( \frac{1}{2} \right),$$

where

- $\widehat{\rho}_{j\ell}(t)$  is the KDE using observations in  $\mathcal{X}_j, \mathcal{X}_\ell$  projected onto the line segment  $\overline{c_j c_\ell}$  and evaluated at  $t \cdot c_j + (1 - t)c_\ell$ .
- Thus,  $\widehat{\rho}_{j\ell} \left( \frac{1}{2} \right)$  is the midpoint (on the boundary).

## Alternatives: Tube density

- Similar to the face density, we also consider the tube density.
- Tube density:

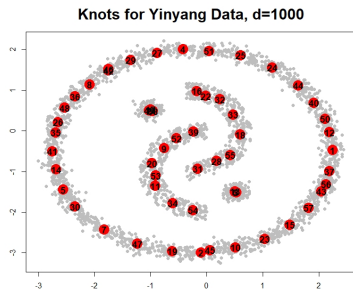
$$\widehat{S}_{j\ell}^{TD} = \min_{t \in [0,1]} \widehat{\omega}_{j\ell,R}(t),$$

where

- $\widehat{\omega}_{j\ell,R}(t)$  is the KDE of all observations projected to  $\overline{c_j c_\ell}$  and evaluated at  $t \cdot c_j + (1 - t)c_\ell$  with a projected distance less than  $R$ .
- Namely,

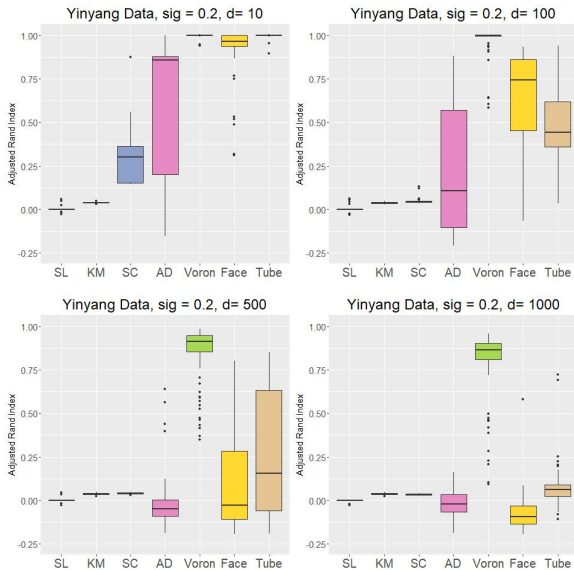
$$\widehat{\omega}_{j\ell,R}(t) = \frac{1}{nh} \sum_{i=1}^n K\left(\frac{\Pi_{j\ell}(X_i) - t \cdot c_j - (1 - t)c_\ell}{h}\right) I(d(X_i, \overline{c_j c_\ell}) < R).$$

# Simulation: Yingyang

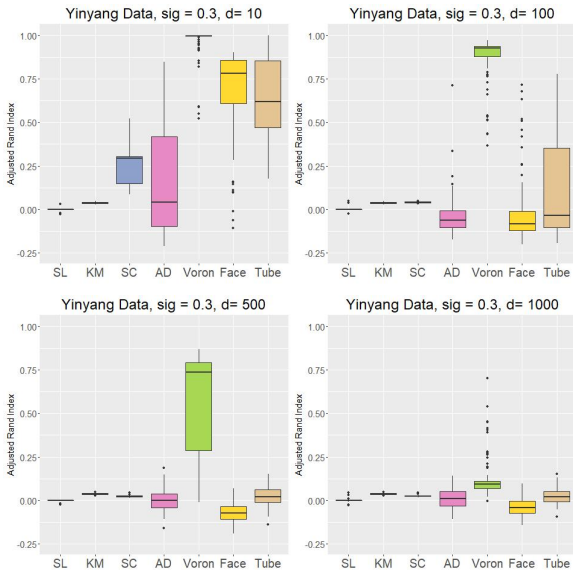


- An yingyang shape data with  $n = 3200$ .
- Main structure is 2D in the region  $[-3, 3] \times [-3, 3]$ .
- We added additional variables to make it high dimensions ( $d = 10, 100, 500, 1000$ ).
- Additional noise level  $\sigma = 0.2, 0.3$ .
- We use adjusted rand index to evaluate performance.

# Simulation: Yinyang ( $\sigma = 0.2$ )

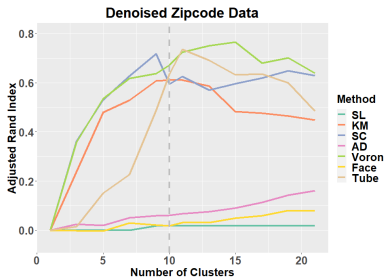
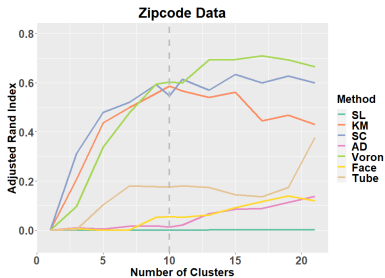


# Simulation: Yingyang ( $\sigma = 0.3$ )



- $n = 2000$  with  $d = 16 \times 16$  images of handwritten Hindu-Arabic numerals from.
- Numbers: 0,1,2,3,4,5,6,7,8,9.
- While this data is often used for classification, we remove the class label and treat it as a clustering problem.
- We consider using the original data and the denoised data (removing observations with the lowest 10% density).

# Real data: Zipcode data - 2



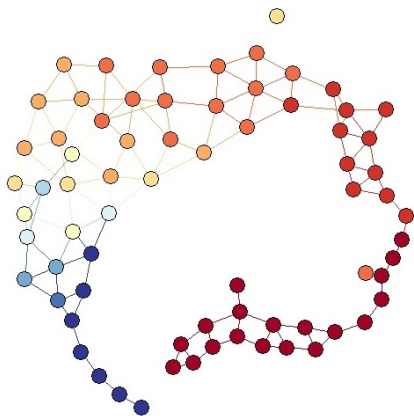
# Skeleton Regression: Introduction

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- The idea of skeleton can be used in a regression setting.
- Intuition: we construct skeleton using the covariates/features and do prediction on the skeleton.

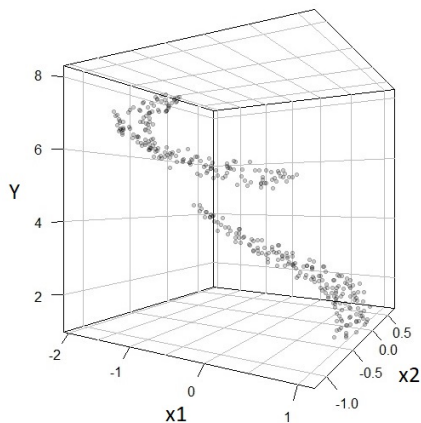


# Skeleton Regression: Application to Astronomy data



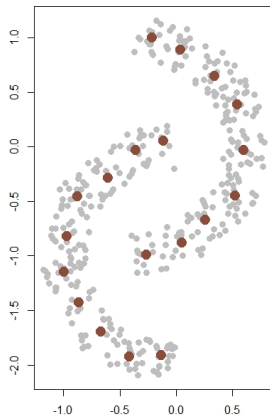
Detecting galaxy's redshift using color information (5D covariates).

# Skeleton Regression: big picture



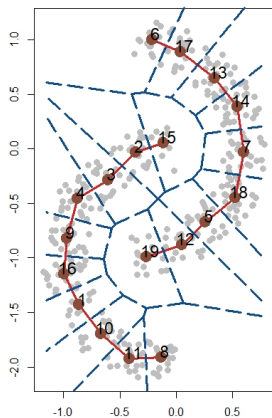
Raw data (2D covariate + 1D response).

# Skeleton Regression: big picture



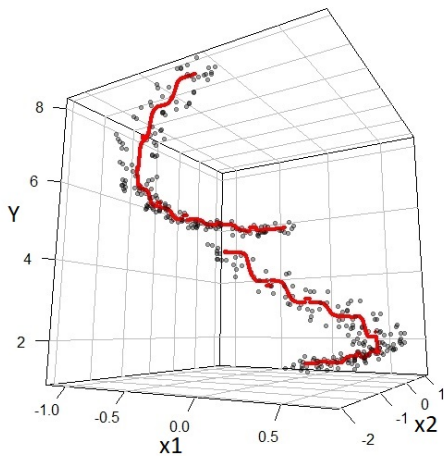
$k$ -means: generating knots (centers of  $k$ -means clusters);  $k = \sqrt{n}$ .

# Skeleton Regression: big picture



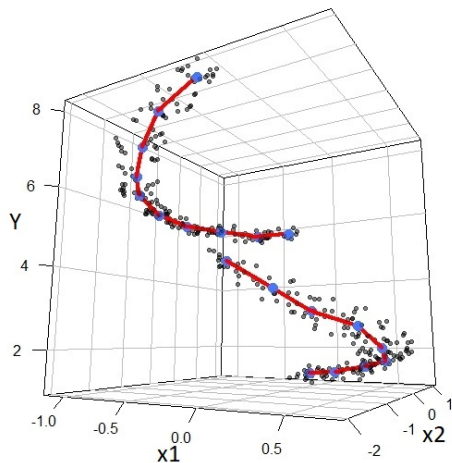
Generating the skeleton.

# Skeleton Regression: big picture



Skeleton kernel regression.

# Skeleton Regression: big picture



Skeleton linear spline.

# Skeleton Regression: Skeleton - 1

- Data:  $(X_1, Y_1), \dots, (X_n, Y_n) \in \mathbb{R}^d \times \mathbb{R}$ .
- We construct skeletons using  $X_1, \dots, X_n$ .
- Centers of k-means:  $c_1, \dots, c_k$ .
- The skeleton clustering creates a weighted graph  $G_0 = (V, W)$ , where
  - $V = \{c_1, \dots, c_k\}$
  - $W = \{w_{j\ell}\}$ , where  $w_{j\ell} = \widehat{S}_{j\ell}^{VD}$  is the Voronoi density.
- We choose a threshold  $\lambda$  to convert it into an unweighted graph  $G = (V, E)$ , where  $e_{j\ell} \in E$  ( $j, \ell$  share an edge) if  $W_{j\ell} \geq \lambda$ .

- The graph  $G$  creates a skeleton  $\mathcal{S} \subset \mathbb{R}^d$  such that

$$\mathcal{S} = V \cup \mathcal{E},$$

where

$$\mathcal{E} = \{tc_j + (1-t)c_\ell : t \in (0,1), e_{j\ell} \in E\}$$

denotes the edges.

- $\mathcal{S}$  is almost a 1D structure except for knots that may have multiple edges attaching to them.
- $\mathcal{S}$  can be decomposed into the vertex region  $V$  and the edge region  $\mathcal{E}$ .
- We project each observation  $X_i$  to  $S_i \in \mathcal{S}$  and construct prediction models accordingly.



- A simple nonparametric regression on skeleton is the linear spline.
- **Skeleton linear spline:** For each point  $s \in \mathcal{S}$ , we require the prediction model  $m(s)$  that
  1.  $m(s)$  is linear when  $s$  is on an edge, and
  2.  $m(s)$  is continuous at each knot.
- While it may look non-trivial to fit this model, there is a simple representer theorem for this.

# Skeleton linear spline: representer theorem

- Define a regression model  $m_\beta$  such that
  - $m_\beta(V_j) = \beta_j$  for each vertex,
  - $m_\beta(s) = t(s)\beta_j + (1 - t(s))\beta_\ell$  if  $s = t(s)V_j + (1 - t(s))V_\ell$ .
- Namely, the model is a linear interpolation of the prediction values on each knot.
- The model  $m_\beta$  is determined by the coefficients  $\beta_1, \dots, \beta_k$  on the knots.

## Theorem (Wei and Chen (2023))

*Any skeleton linear spline model can be written as  $m_\beta$  for some  $\beta$ .*

## Skeleton linear spline: fitting

- Fitting the skeleton linear spline is very easy.
- For every observation  $X_i$  with a projected location  $S_i \in \mathcal{S}$ , we further convert it into a vector  $Z_i \in [0, 1]^k$  such that

$$Z_{ik} = \begin{cases} 1 & \text{if } S_i = V_k \text{ is on the vertex} \\ t & \text{if } S_i = tV_k + (1-t)V_\ell \text{ for some } V_\ell \\ 0 & \text{otherwise.} \end{cases}$$

- With this, the prediction value  $m_\beta(S_i) = Z_i^T \beta$ .
- Thus, when we estimate  $\beta$  using the least square, this becomes a linear regression problem with an analytic solution:

$$\hat{\beta} = (ZZ^T)^{-1}ZY.$$

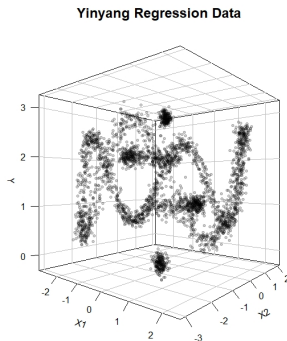
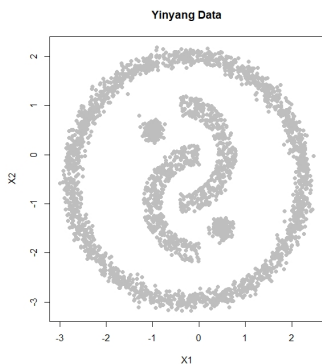
- The set  $\mathcal{S}$  is equipped with a metric  $d_{\mathcal{S}}$  because
  - each vertex  $c_j \in \mathbb{R}^d$  has a location in Euclidean space and
  - each edge  $e_{j\ell}$  has a length  $\|c_j - c_{\ell}\|$ .
- For two points  $s_1, s_2 \in \mathcal{S}$ , their distance  $d_{\mathcal{S}}(s_1, s_2)$  will be the shortest distance in  $\mathcal{S}$ . If they belong to two different connected component, we set  $d_{\mathcal{S}}(s_1, s_2) = \infty$ .
- The metric space  $(\mathcal{S}, d_{\mathcal{S}})$  allows us to use a wide variety of methods for prediction.

- As a classical example, we may use kernel regression on the skeleton.
- The prediction value  $\hat{m}_h(s)$  is

$$\hat{m}_h(s) = \frac{\sum_{i=1}^n Y_i K\left(\frac{d_S(s, S_i)}{h}\right)}{\sum_{j=1}^n K\left(\frac{d_S(s, S_j)}{h}\right)}.$$

- Other methods such as kNN is applicable as well.

# Skeleton Regression: simulations - 1



We add additional covariates to make it a high-dimensional data.

## Skeleton Regression: simulations - 2

Method	Medium SSE (5%, 95%)	nknots	Parameter
kNN	204.5 (192.3, 221.9)	-	neighbor=18
Ridge	2127.0 (2100.2, 2155.2)		$\lambda = 7.94$
Lasso	1556.8 (1515.4, 1607.9)		$\lambda = 0.0126$
SpecSeries	1506.4 (1469.1, 1555.6)	-	bandwidth = 2
S-Kernel	112.8 (102.0, 121.7)	38	bandwidth = 6 $r_{hns}$
S-kNN	139.6 (129.6, 148.7)	38	neighbor = 36
S-Lspline	95.8 (88.6, 102.6)	38	-

$d = 1000$ . We use 10-fold cross-validation for every method.

- Skeleton approach offers a flexible framework.
- It shows promising results in both clustering and regression when the number of covariates is high.
- However, a couple of open questions remains:
  - Understanding the effect of  $k$ -means when  $k$  is large.
  - How does the randomness of  $k$ -means affects the final result.
  - Principled way to post-process the knots.
- Main references:
  - Skeleton clustering: arXiv 2104.10770
  - Skeleton regression: arXiv 2303.11786



# Thank You!

More details can be found in  
<http://faculty.washington.edu/yenchic>.