### Bones, Teeth and Animation

# **Ingrid Daubechies**

### May 10, 2016

### Green Family Lecture, IPAM, UCLA

surfaces and Morphology

### Collaborators



Rima Alaifari ETH Zürich



Doug Boyer Duke



Yaron Lipman Weizmann



Roi Poranne ETH Zürich



Ingrid Daubechies Duke



Jesús Puente J.P. Morgan



Tingran Gao Duke



Robert Ravier Duke

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・

₹ 9 Q

Machine Learning, Fibre Bundles and Biological Morphology

Ingrid Daubechies Tingran Gao

Department of Mathematics Duke University

Feb 11, 2016

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > <

• 3-dimensional Animation relies on computer graphics

- 3-dimensional Animation relies on computer graphics
- computer graphics uses 3-dimensional mesh models

In the beginning was the TEAPOT

### TEAPOT:



Distances between Surfaces

ヘロト 人間 とくほ とくほとう

ъ

### TEAPOT:



Distances between Surfaces

3D meshes became much more sophisticated over the years

◆□▶ ◆□▶ ◆ 三▶ ◆ 三▶ - 三 - のへぐ



#### Distances between Surfaces

<ロ> <四> <四> <三</p>





<ロ> <同> <同> <同> < 同> < 同>

ъ

DQC



Distances between Surfaces

<ロ> <四> <四> <四> <三</td>





Distances between Surfaces

<ロ> <四> <四> <三</p>

Scan existing objects

Distances between Surfaces

◆ロト ◆御 ト ◆臣 ト ◆臣 ト ─ 臣 … のへで

### Scan existing objects



#### Distances between Surfaces

<ロ> <四> <四> <四> <三</td>

### Scan existing objects



### Scan existing objects





イロト イポト イヨト イヨト

Э

DQC

- point cloud → triangulation (Delauney triangulation)
- ۲
- ٩
- •





イロト イポト イヨト イヨト

Э

Dac

- point cloud → triangulation (Delauney triangulation)
- ۲
- ٩
- •
- •



◆ロ → ◆檀 → ◆注 → ◆注 → □ 注

DQC

- point cloud → triangulation (Delauney triangulation)
- edit triangulated surfaces
- ٩
- ٠
- •



イロト 不得 トイヨト イヨト

Э

- point cloud → triangulation (Delauney triangulation)
- edit triangulated surfaces
- recognize identical surfaces?
- ۹
- ۹



ヘロア 人間 アメヨア 人口 ア

Э

- point cloud → triangulation (Delauney triangulation)
- edit triangulated surfaces
- recognize identical surfaces?
- or deformations of each other?



ヘロア 人間 アメヨア 人口 ア

Э

- point cloud → triangulation (Delauney triangulation)
- edit triangulated surfaces
- recognize identical surfaces?
- or deformations of each other?
- quantify difference?





・ロト ・ 同ト ・ ヨト ・ ヨト

# **Reference points**

### Animating "humanoid" characters requires reference points



Distances between Surfaces

イロト イボト イヨト イヨト 二日

# **Reference points**

Animating "humanoid" characters requires reference points



イロト 不得 トイヨト イヨト

3

# **Reference points**

### Animating "humanoid" characters requires reference points



Distances between Surfaces

<ロ> <四> <四> <四> <三</td>

We need to be able to:

- recognize when two point clouds correspond to the same surface or to two similar surfaces
- quantify how different two surfaces are from each other (or how similar to each other)
- find correspondence points for similar surfaces

<ロト < 同ト < 巨ト < 巨ト < 巨 > つへの

## It all started with a conversation with biologists....





#### Jukka Jernvall

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > <

More Precisely: biological morphologists Study Teeth & Bones of extant & extinct animals still live today fossils First: project on "complexity" of teeth

◆□▶ ◆□▶ ◆臣▶ ◆臣▶ 臣 の�?



Landmarked Teeth 
$$\longrightarrow$$
  
 $d_{Procrustes}^{2}\left(S_{1}, S_{2}\right) = \min_{R \text{ rigid tr.}} \sum_{j=1}^{J} \left\|R\left(x_{j}\right) - y_{j}\right\|^{2}$ 









Landmarked Teeth 
$$\longrightarrow$$
  
 $d_{Procrustes}^{2}\left(S_{1}, S_{2}\right) = \min_{R \text{ rigid tr.}} \sum_{j=1}^{J} \left\|R\left(x_{j}\right) - y_{j}\right\|^{2}$ 



Find way to compute a distance that does as well, for biological purposes, as Procrustes distance, based on expert-placed landmarks, automatically?







Landmarked Teeth 
$$\longrightarrow$$
  
 $d_{Procrustes}^{2}\left(S_{1}, S_{2}\right) = \min_{R \text{ rigid tr.}} \sum_{j=1}^{J} \left\|R\left(x_{j}\right) - y_{j}\right\|^{2}$ 



Find way to compute a distance that does as well, for biological purposes, as Procrustes distance, based on expert-placed landmarks, automatically?

examples: finely discretized triangulated surfaces









# We defined 2 different distances

- $d_{
  m cWn}(S_1,S_2)$ : conformal flattening comparison of neighborhood geometry optimal mass transport
  - $d_{\mathrm{cP}}\left(S_{1},S_{2}
    ight)$ : continuous Procrustes distance











Even mistake made by  $d_{\rm cP}$ were similar to biologists' mistakes

small distances between  $S_1, S_2 \longrightarrow OK$  maps larger distances  $\longrightarrow$  not OK

## Biologists' "wish list" changed...

- ... as they learned our language and saw our methods
  - mappings more important to them than distances
     (---> discussion of variability in individuals or between species, locally)

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > <

no holonomy!
## Biologists' "wish list" changed...

- ... as they learned our language and saw our methods
  - mappings more important to them than distances
    (--> discussion of variability in individuals or between species, locally)
  - no holonomy!

### Our formulation of problem changed too

 $\begin{array}{rcl} \mbox{Tingran Gao} & \longrightarrow & \mbox{reformulate as connection on fibre bundle} \\ & + & \mbox{horizontal diffusion} \end{array}$ 

< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > <

# Even before this...

biological content in large concatenated matrix



◆□▶ ◆□▶ ◆三▶ ◆三▶ 三三 のへぐ

# Even before this...

biological content in large concatenated matrix



$$\begin{array}{l} \min \ \left\|\boldsymbol{M}\mathbb{X}-\mathbb{X}\right\|_{2}^{2}+\lambda \left\|\mathbb{X}\right\|_{1}\\ \text{s.t.} \ \left\|\mathbb{X}\right\|_{2}=1. \end{array}$$

Resulting minimizers X supported on <u>union</u> of 4 surfaces

Use the Information in the Maps!

$$d_{\mathrm{cP}}\left(S_{1},S_{2}\right) = \inf_{\mathcal{C}\in\mathcal{A}\left(S_{1},S_{2}\right)} \inf_{R\in\mathbb{E}(3)} \left(\int_{S_{1}} \left\|R\left(x\right)-\mathcal{C}\left(x\right)\right\|^{2} d\mathrm{vol}_{S_{1}}\left(x\right)\right)^{\frac{1}{2}}$$



・ロト ・ 日 ト ・ モ ト ・ モ ト

## Learning from Distances



・ロト ・聞ト ・ヨト ・ヨト



## Learning from Distances



◆□> ◆□> ◆豆> ◆豆> ・豆 ・ のへで

## Learning from Distances



#### **Diffusion Distance**

・ロト ・聞ト ・ヨト ・ヨト

- 2

# MDS for CPD & DD





CPD

DD

・ロト ・聞ト ・ヨト ・ヨト

# MDS for CPD & DD





CPD

DD

・ロト ・聞ト ・ヨト ・ヨト



◆ロ ▶ ◆母 ▶ ◆臣 ▶ ◆臣 ▶ ● ● ● ● ●



▲□▶ ▲圖▶ ▲厘▶ ▲厘▶ 厘 の��



◆□▶ ◆□▶ ◆臣▶ ◆臣▶ ─臣 ─のへで



◆□▶ ◆□▶ ◆臣▶ ◆臣▶ ─臣 ─のへで



◆□▶ ◆□▶ ◆臣▶ ◆臣▶ ─臣 ─ のへで



◆□ > ◆□ > ◆豆 > ◆豆 > ̄豆 \_ のへぐ



・ロ・・(型・・(目・・(目・・)のへぐ



•  $P = D^{-1}W$  defines a random walk on the graph

▲□▶ ▲圖▶ ★ 国▶ ★ 国▶ - 国 - のへで



- $P = D^{-1}W$  defines a random walk on the graph
- Solve eigen-problem

$$Pu_j = \lambda_j u_j, \ j = 1, 2, \cdots, m$$

▲□▶ ▲圖▶ ★ 国▶ ★ 国▶ - 国 - のへで



- $P = D^{-1}W$  defines a random walk on the graph
- Solve eigen-problem

$$Pu_j = \lambda_j u_j, \ j = 1, 2, \cdots, m$$

and represent each individual shape  $S_i$  as an *m*-vector

$$\left(\lambda_{1}^{t/2}u_{1}\left(j\right),\cdots,\lambda_{m}^{t/2}u_{m}\left(j\right)\right)$$

◆□▶ ◆□▶ ◆三▶ ◆三▶ 三三 のへぐ

# Diffusion Distance (DD) Fix $1 \le m \le N$ , $t \ge 0$ ,

$$D_{m}^{t}(S_{i}, S_{j}) = \left(\sum_{k=1}^{m} \lambda_{k}^{t} (u_{k}(i) - u_{k}(j))^{2}\right)^{\frac{1}{2}}$$

◆□▶ ◆□▶ ◆臣▶ ◆臣▶ 三臣 - のへで

# Diffusion Distance (DD) Fix $1 \le m \le N$ , $t \ge 0$ ,

$$D_m^t(S_i, S_j) = \left(\sum_{k=1}^m \lambda_k^t \left(u_k(i) - u_k(j)\right)^2\right)^{\frac{1}{2}}$$



▲□▶ ▲圖▶ ▲臣▶ ▲臣▶ 三臣 - のへで

# MDS for CPD & DD





CPD

DD

・ロト ・聞ト ・ヨト ・ヨト

# Even Better: More Information!





HBDD

DD

<ロト <回ト < 注ト < 注ト

# Even Better: More Information!





HBDD

DD

<ロト <回ト < 注ト < 注ト

Use the Information in the Maps!

$$D\left(S_{1},S_{2}\right) = \inf_{\mathcal{C}\in\mathcal{A}\left(S_{1},S_{2}\right)} \inf_{R\in\mathbb{E}\left(3\right)} \left(\int_{S_{1}} \|R\left(x\right) - \mathcal{C}\left(x\right)\|^{2} d\operatorname{vol}_{S_{1}}\left(x\right)\right)^{\frac{1}{2}}$$



・ロト ・ 日 ト ・ モ ト ・ モ ト



◆□ > ◆□ > ◆豆 > ◆豆 > ̄豆 \_ のへぐ



▲口▶ ▲圖▶ ▲臣▶ ▲臣▶ 三臣 - 釣�??



э

< ロ > < 同 > < 三 > <



▲口▼▲□▼▲目▼▲目▼ 目 めんぐ



▲ □ ▶ ▲ 圖 ▶ ▲ 圖 ▶ ▲ 圖 ■ め Q @ .

# Distance Graph



◆□▶ ◆□▶ ◆臣▶ ◆臣▶ 臣 の�?

# Augmented Distance Graph



◆□▶ ◆□▶ ◆ □▶ ◆ □▶ ○ □ ○ ○ ○ ○

Fibre Bundle  $\mathscr{E} = (E, M, F, \pi)$ 

- E: total manifold
- M: base manifold
- $\pi: E \to M$ : smooth surjective map (bundle projection)

▲ロト ▲帰 ト ▲ ヨ ト ▲ ヨ ト ・ ヨ ・ の Q ()

F: fibre manifold

Fibre Bundle  $\mathscr{E} = (E, M, F, \pi)$ 

- E: total manifold
- M: base manifold
- $\pi: E \to M$ : smooth surjective map (bundle projection)
- F: fibre manifold
- Iocal triviality: for "small" open set U ⊂ M, π<sup>-1</sup>(U) is diffeomorphic to U × F





Fibre Bundle  $\mathscr{E} = (E, M, F, \pi)$ 

- E: total manifold
- M: base manifold
- $\pi: E \to M$ : smooth surjective map (bundle projection)
- F: fibre manifold
- Iocal triviality: for "small" open set U ⊂ M, π<sup>-1</sup>(U) is diffeomorphic to U × F



Fibre Bundle  $\mathscr{E} = (E, M, F, \pi)$ 

- E: total manifold
- M: base manifold
- $\pi: E \to M$ : smooth surjective map (bundle projection)
- F: fibre manifold
- Iocal triviality: for "small" open set U ⊂ M, π<sup>-1</sup>(U) is diffeomorphic to U × F


- E: total manifold
- M: base manifold
- $\pi: E \to M$ : smooth surjective map (bundle projection)
- F: fibre manifold
- Iocal triviality: for "small" open set U ⊂ M, π<sup>-1</sup>(U) is diffeomorphic to U × F



- E: total manifold
- M: base manifold
- $\pi: E \to M$ : smooth surjective map (bundle projection)
- F: fibre manifold
- Iocal triviality: for "small" open set U ⊂ M, π<sup>-1</sup>(U) is diffeomorphic to U × F



- E: total manifold
- M: base manifold
- $\pi: E \to M$ : smooth surjective map (bundle projection)
- F: fibre manifold
- Iocal triviality: for "small" open set U ⊂ M, π<sup>-1</sup>(U) is diffeomorphic to U × F



- E: total manifold
- M: base manifold
- $\pi: E \to M$ : smooth surjective map (bundle projection)
- F: fibre manifold
- Iocal triviality: for "small" open set U ⊂ M, π<sup>-1</sup>(U) is diffeomorphic to U × F



Diffusion Maps

$$D^{-1}Wu_k = \lambda_k u_k, \quad 1 \le k \le N$$



◆□▶ ◆□▶ ◆臣▶ ◆臣▶ = 臣 = のへで

Horizontal Diffusion Maps

$$\mathcal{D}^{-1}\mathcal{W}u_k = \lambda_k u_k, \quad 1 \le k \le \kappa$$



・ロット (日) (日) (日) (日) (日)

Horizontal Diffusion Maps

$$\mathcal{D}^{-1}\mathcal{W}u_k = \lambda_k u_k, \quad 1 \le k \le \kappa$$



・ロット (日) (日) (日) (日) (日)

Horizontal Diffusion Maps

$$\mathcal{D}^{-1}\mathcal{W}u_{k} = \lambda_{k}u_{k}, \quad 1 \leq k \leq \kappa$$

$$\mathcal{D}^{-1}\begin{pmatrix} & \vdots & \\ & \vdots & \\ & & e^{-d_{ij}^{2}/\epsilon}\rho_{ij}^{\delta} & \cdots \\ & & \vdots & \end{pmatrix}\begin{pmatrix} \vdots & \\ \vdots & \\ u_{k[j]} \\ \vdots \end{pmatrix} = \lambda_{k}\begin{pmatrix} \vdots & \\ \vdots \\ u_{k[j]} \\ \vdots \end{pmatrix}$$

Horizontal Diffusion Maps: For fixed  $1 \le m \le \kappa$ ,  $t \ge 0$ , represent  $S_j$  as a  $\kappa_j \times m$  matrix

$$\left(\lambda_1^{t/2}u_{1[j]},\cdots,\lambda_m^{t/2}u_{m[j]}\right)$$

#### Diffusion Maps vs. Horizontal Diffusion Maps

Diffusion Maps: For fixed  $1 \le m \le \kappa$ ,  $t \ge 0$ , represent  $S_j$  as an *m*-dimensional vector

$$\left(\lambda_1^{t/2}u_1(j),\cdots,\lambda_m^{t/2}u_m(j)\right)$$

Horizontal Diffusion Maps: For fixed  $1 \le m \le \kappa$ ,  $t \ge 0$ , represent  $S_j$  as a  $\kappa_j \times m$  matrix

$$\left(\lambda_1^{t/2}u_{1[j]},\cdots,\lambda_m^{t/2}u_{m[j]}\right)$$

### HDM: Application in Geometric Morphometrics

◆□▶ ◆□▶ ◆三▶ ◆三▶ 三三 のへぐ

- 1. Global Registration
- 2. Automatic Landmarking
- 3. Species Classification





▲□▶ ▲□▶ ▲臣▶ ▲臣▶ 三臣 - のへの



▲□▶ ▲□▶ ▲三▶ ▲三▶ 三三 のへの



▲□▶ ▲圖▶ ▲≣▶ ▲≣▶ = 差 = のへで



▲□> ▲圖> ▲目> ▲目> 二目 - のへで





- \* ロ \* \* 個 \* \* 目 \* \* 目 \* \* の < ??

### 2. Automatic Landmarking: Spectral Clustering



### 2. Automatic Landmarking: Spectral Clustering



Horizontal Diffusion Maps (HDM): For fixed  $1 \le m \le \kappa$ ,  $t \ge 0$ , represent  $S_i$  as a  $\kappa_i \times m$  matrix

$$\left(\lambda_1^{t/2}u_{1[j]},\cdots,\lambda_m^{t/2}u_{m[j]}\right)$$

Horizontal Diffusion Maps (HDM): For fixed  $1 \le m \le \kappa$ ,  $t \ge 0$ , represent  $S_j$  as a  $\kappa_j \times m$  matrix

$$\left(\lambda_1^{t/2}u_{1[j]},\cdots,\lambda_m^{t/2}u_{m[j]}\right)$$

Horizontal Base Diffusion Maps (HBDM): For fixed  $1 \le m \le \kappa, t \ge 0$ , represent  $S_j$  as a  $\binom{m}{2}$ -dimensional vector  $\left(\lambda_{\ell}^{t/2}\lambda_{k}^{t/2} \langle u_{\ell[j]}, u_{k[j]} \rangle\right)_{1 \le \ell < k \le m}$ 

Horizontal Base Diffusion Distance (HBDD): For fixed  

$$1 \le m \le \kappa, t \ge 0,$$
  
 $D_{HB}^{t}(S_{i}, S_{j}) = \left(\sum_{1 \le \ell < k \le m} \lambda_{\ell}^{t} \lambda_{k}^{t} \left( \langle u_{\ell[i]}, u_{k[i]} \rangle - \langle u_{\ell[j]}, u_{k[j]} \rangle \right)^{2} \right)^{\frac{1}{2}}$ 

Horizontal Base Diffusion Maps (HBDM): For fixed  $1 \le m \le \kappa, t \ge 0$ , represent  $S_j$  as a  $\binom{m}{2}$ -dimensional vector  $\left(\lambda_{\ell}^{t/2}\lambda_k^{t/2} \langle u_{\ell[j]}, u_{k[j]} \rangle\right)_{1 \le \ell < k \le m}$ 





HBDD

DD

・ロト ・ 日 ・ ・ ヨ ・

æ

э





HBDD

DD

・ロト ・ 日 ・ ・ ヨ ・

æ

э





◆□▶ ◆□▶ ◆臣▶ ◆臣▶ ─臣 ─ のへで



◆□▶ ◆□▶ ◆臣▶ ◆臣▶ 三臣 - のへで