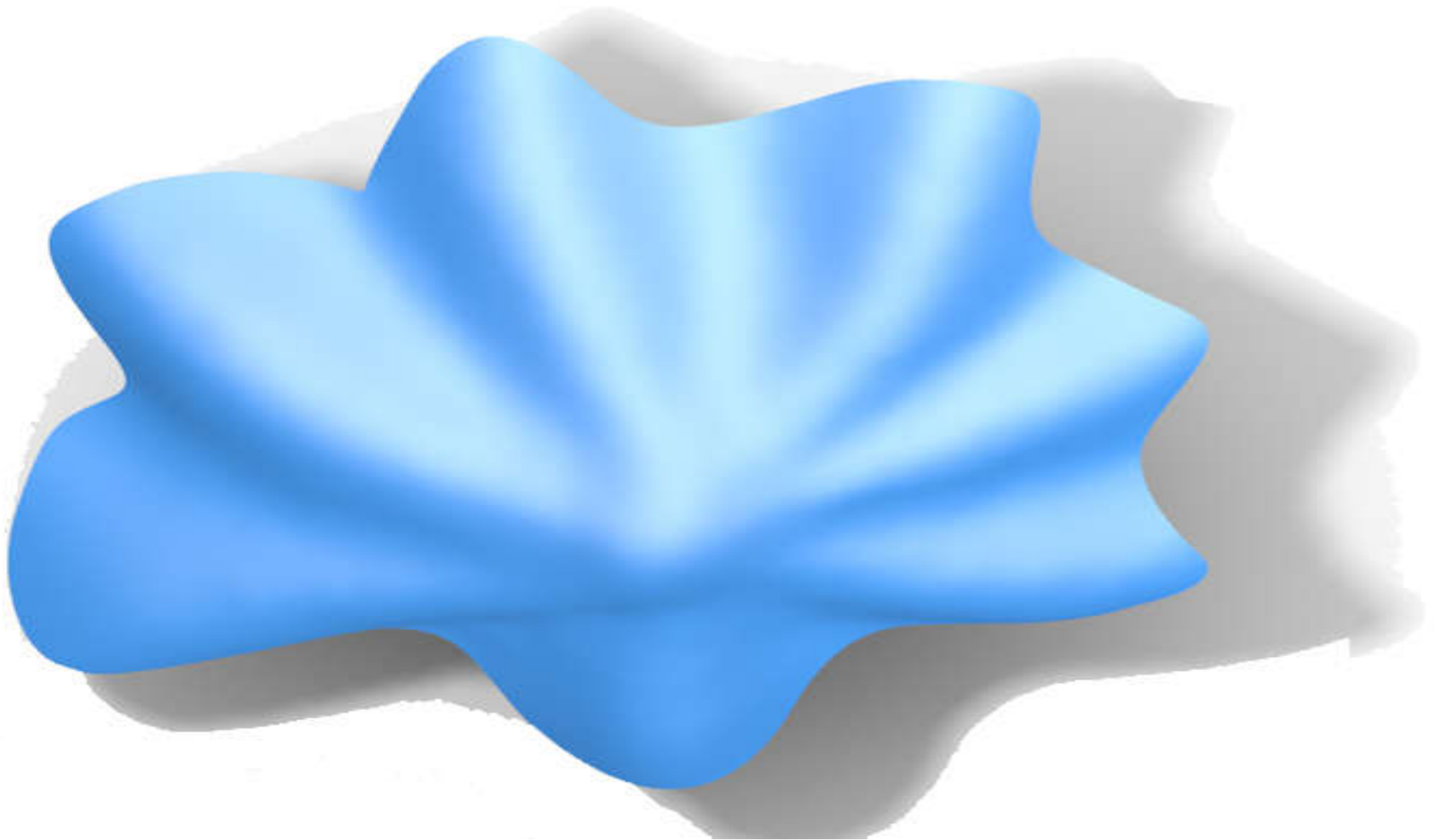


Discovering low-dimensional manifolds in high-dimensional data

Ingrid Daubechies
Duke University

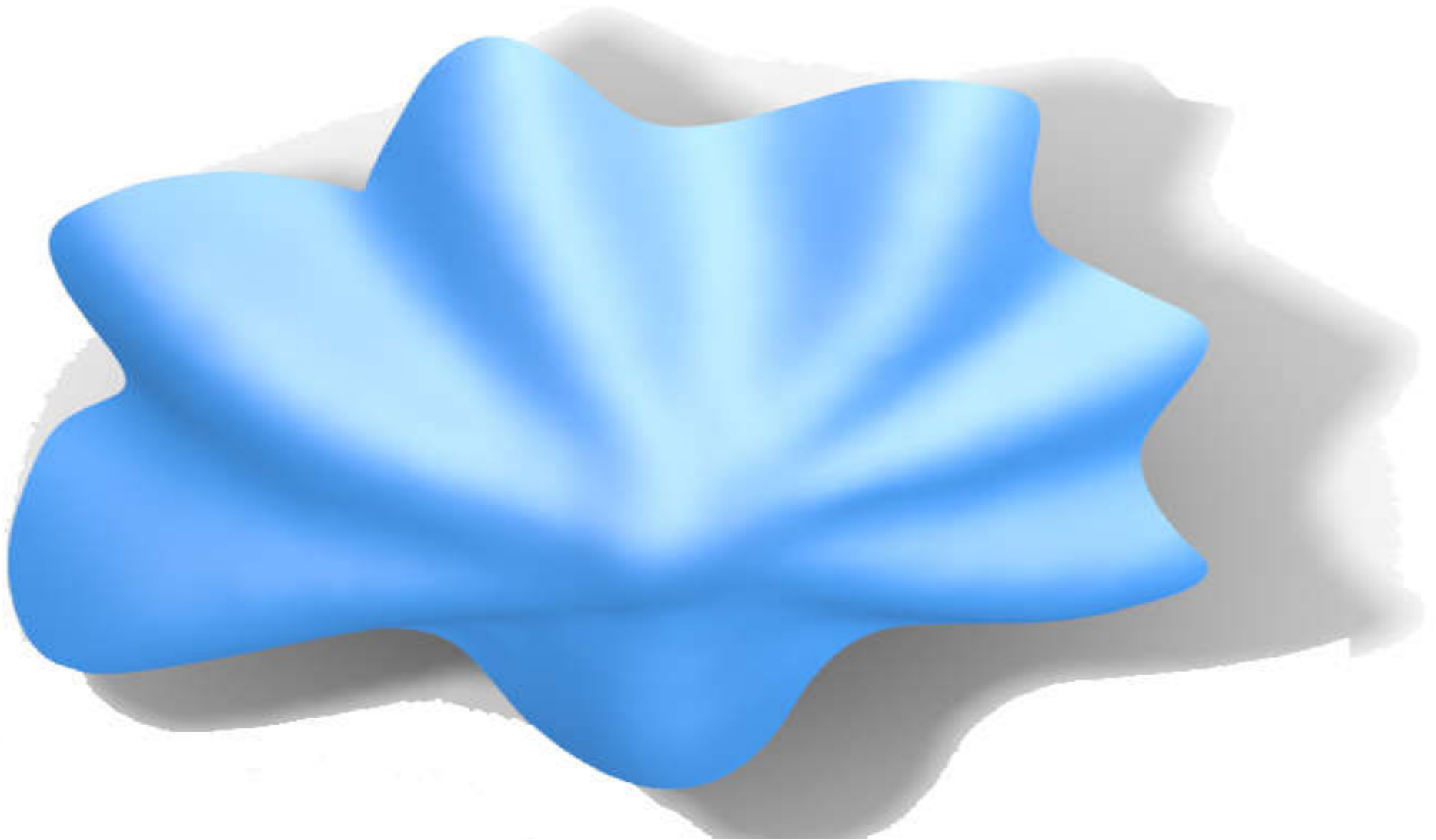
Paris, July 4, 2025

A manifold or variety may have a smaller dimension than how it is presented at first



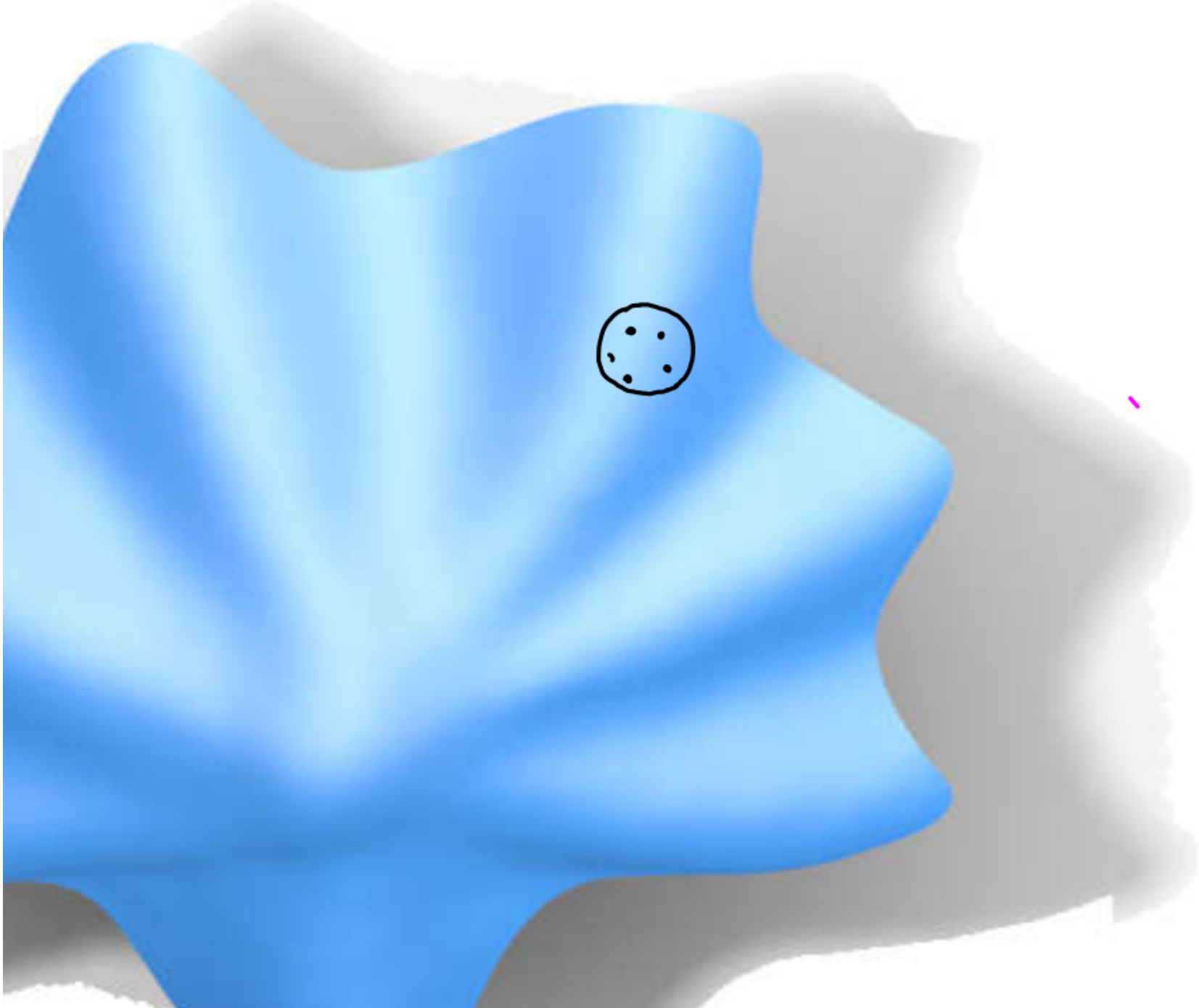
Sample points: given by **3** coordinates (x,y,z)

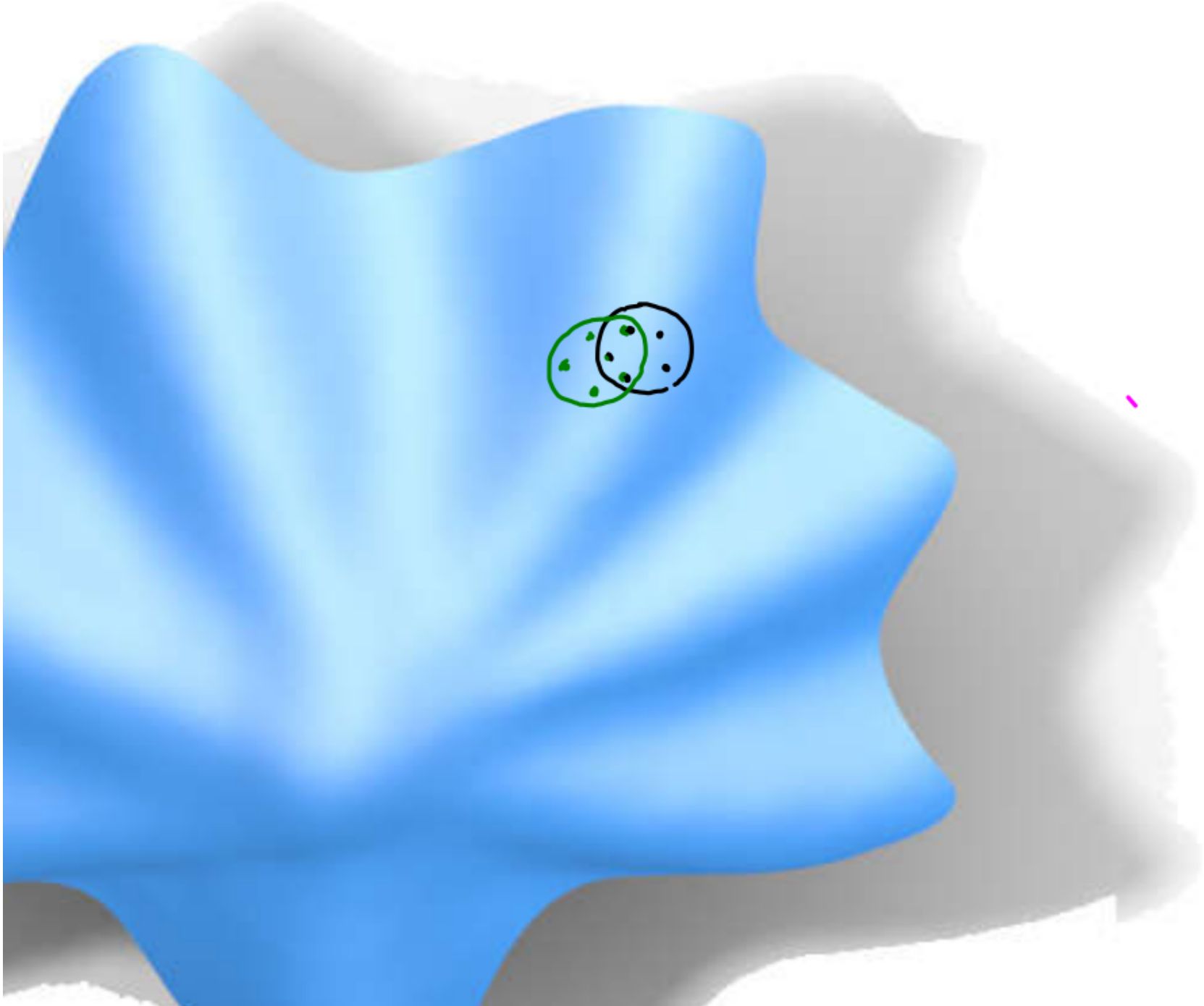
but the blue surface is only **2**-dimensional

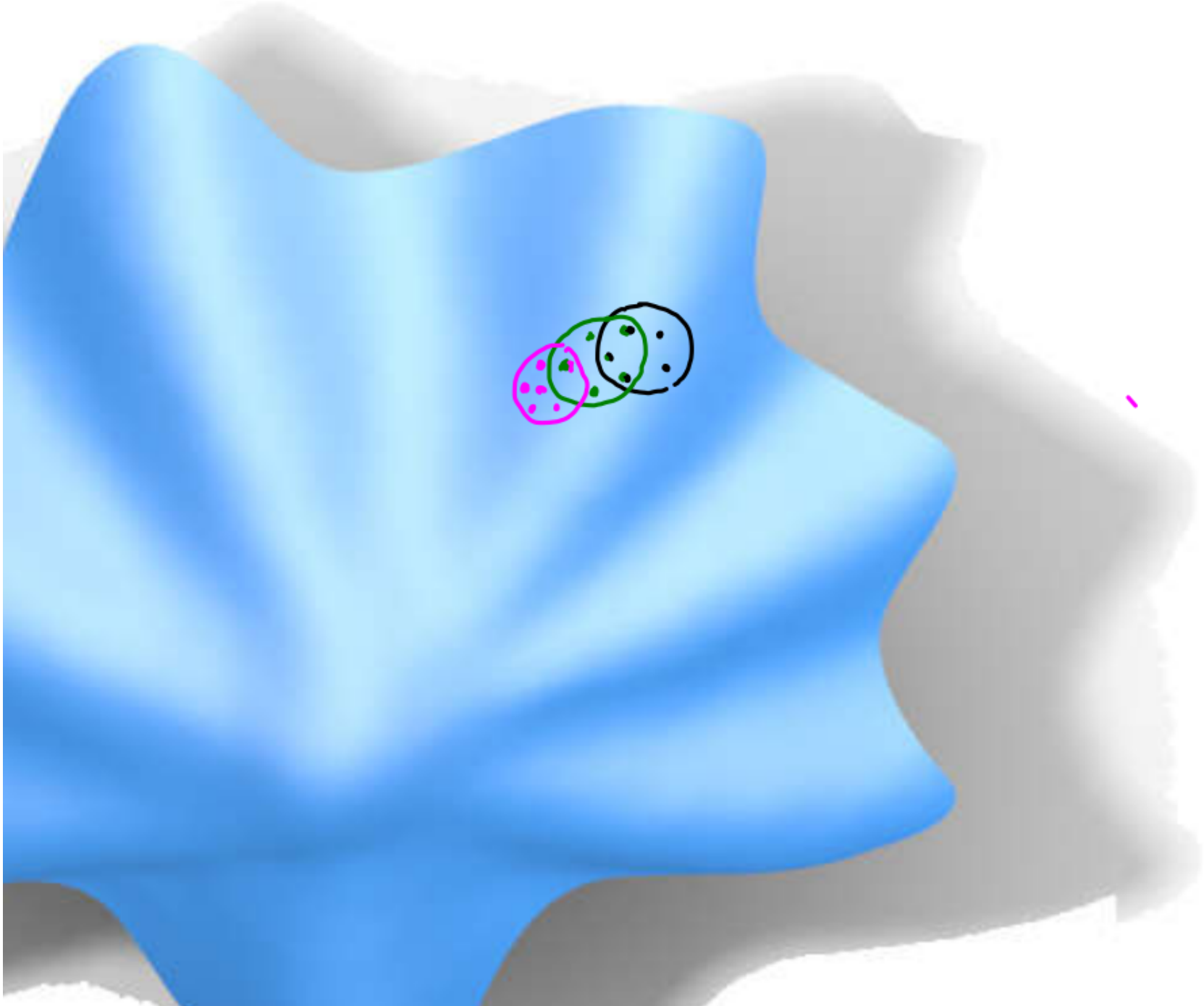


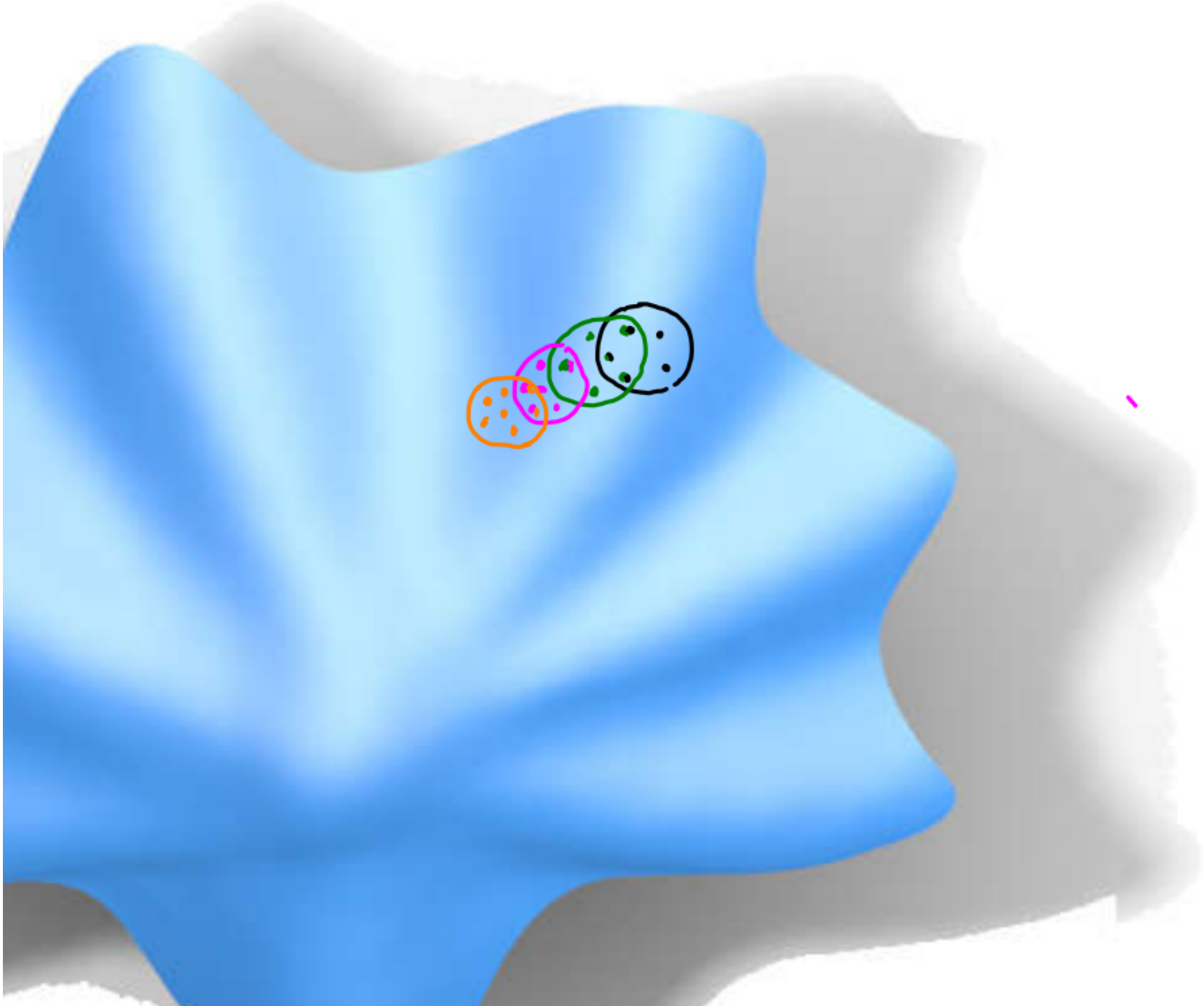
A collection of small tangent patches does give a good first approximation of a surface

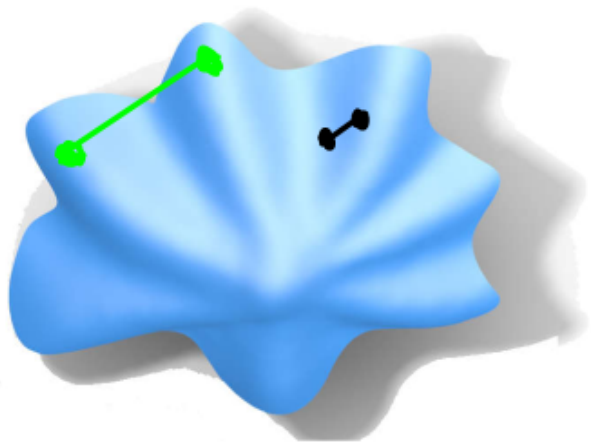




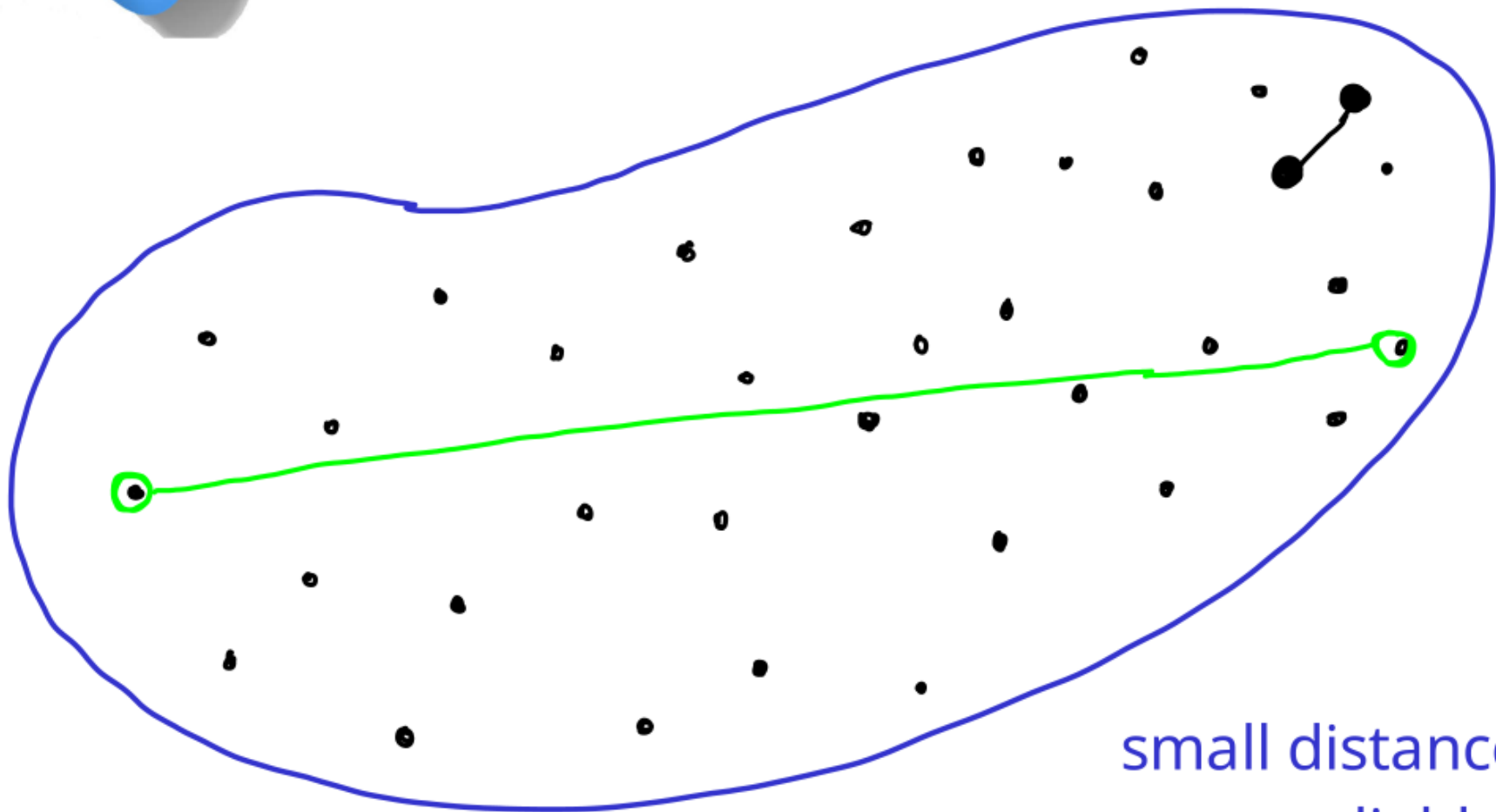




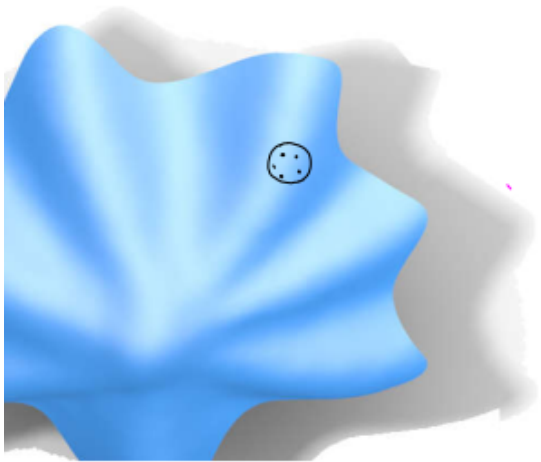




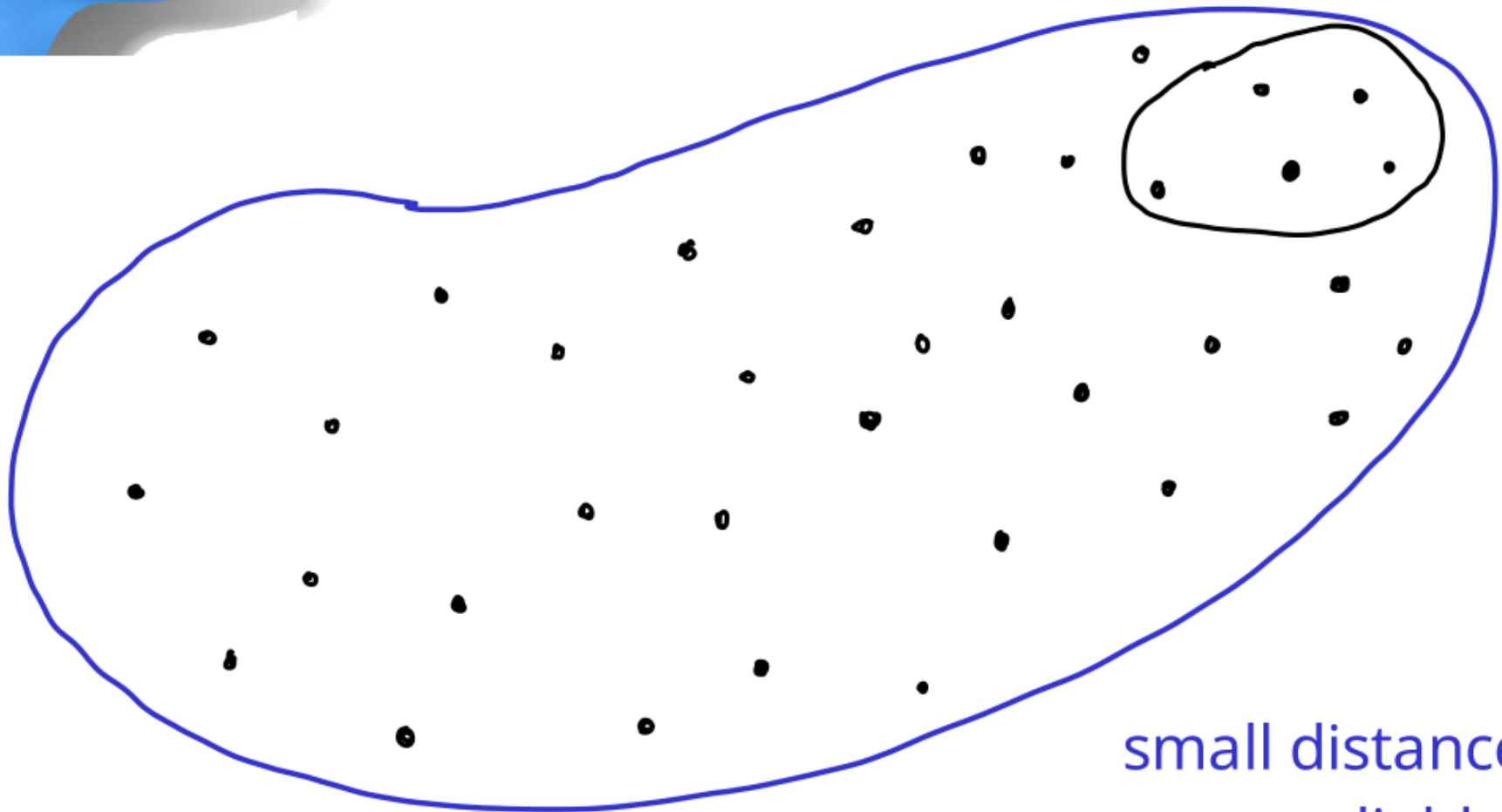
Data set



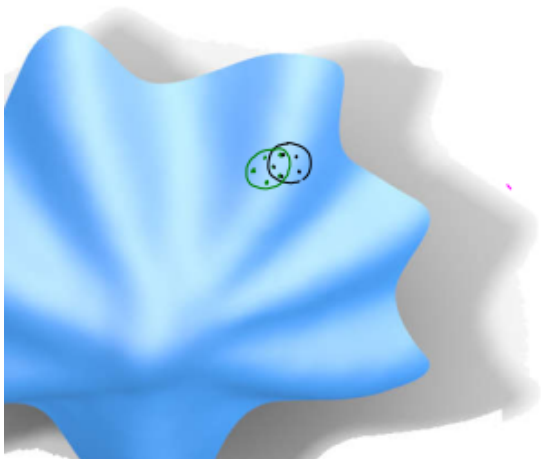
small distances are
more reliable!



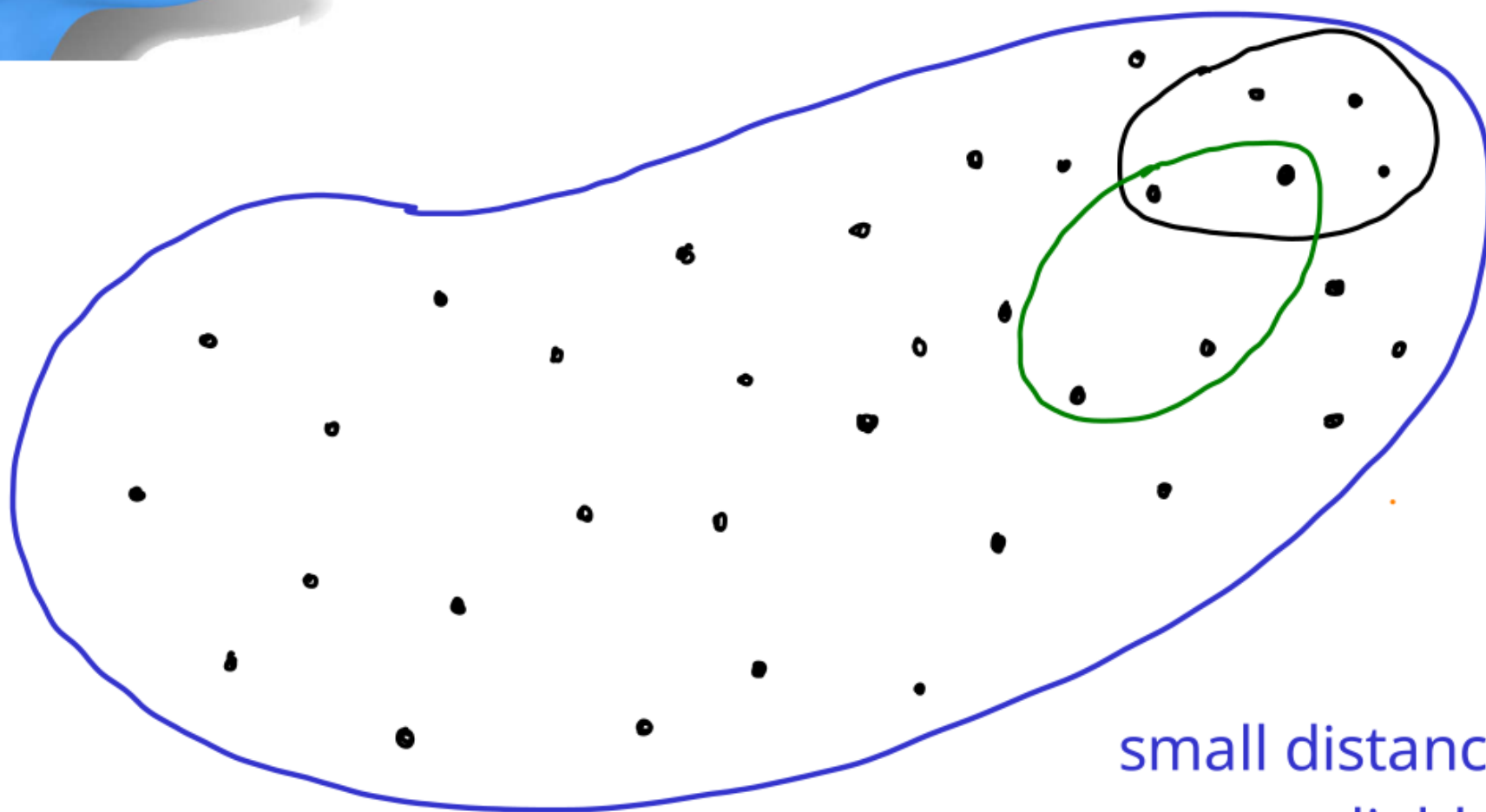
Data set



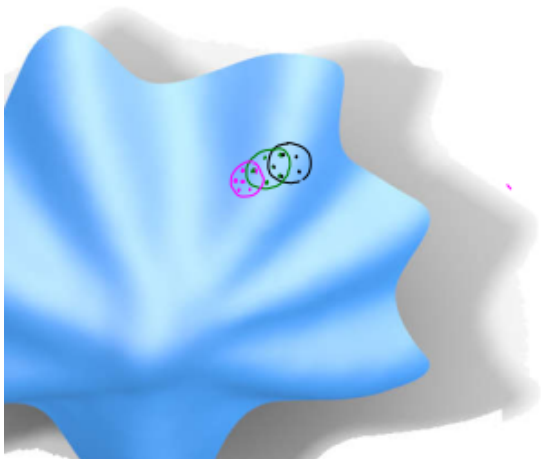
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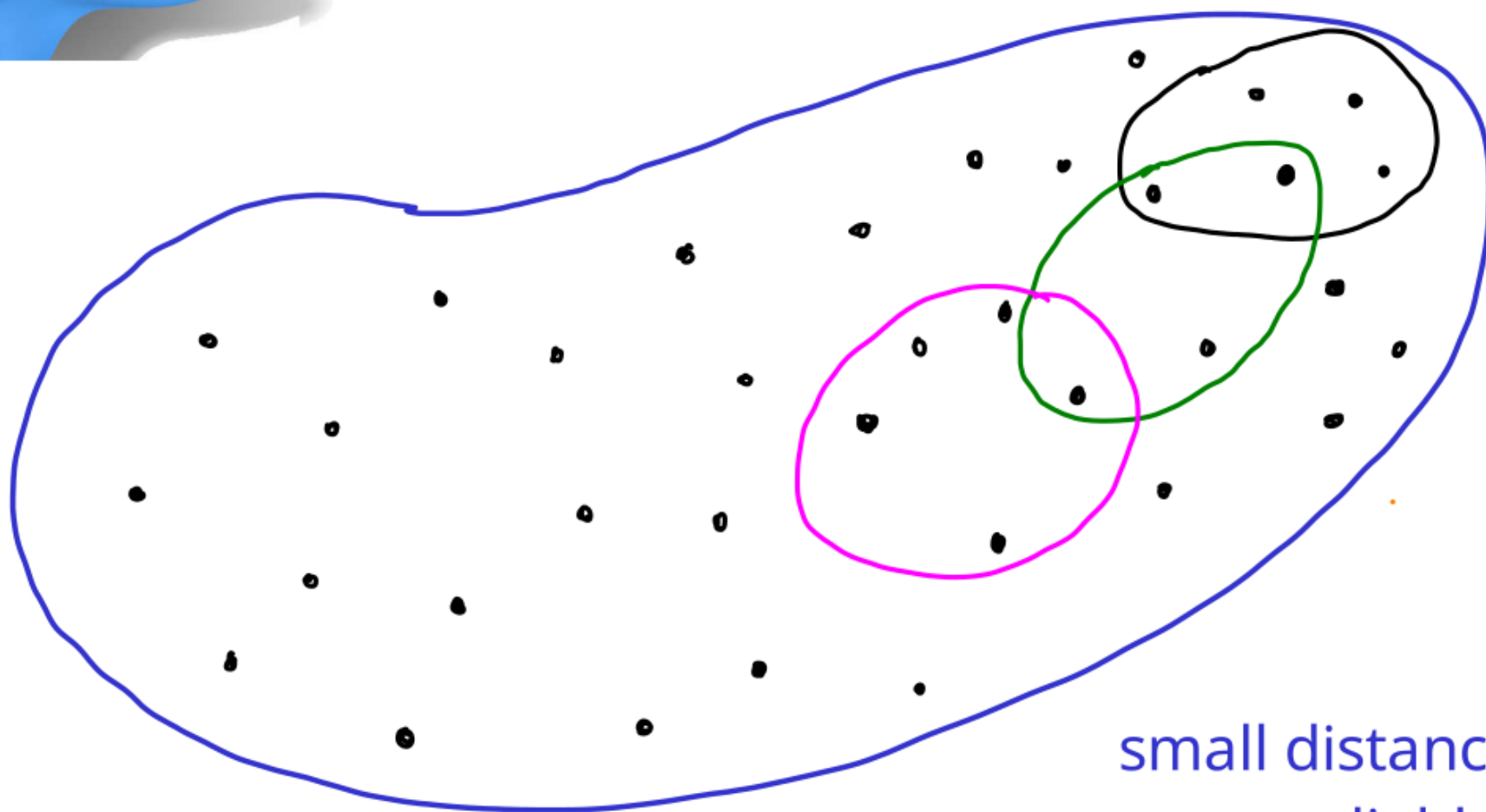
Data set



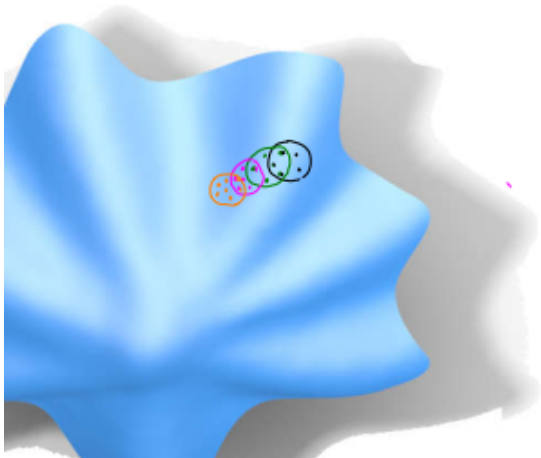
small distances are
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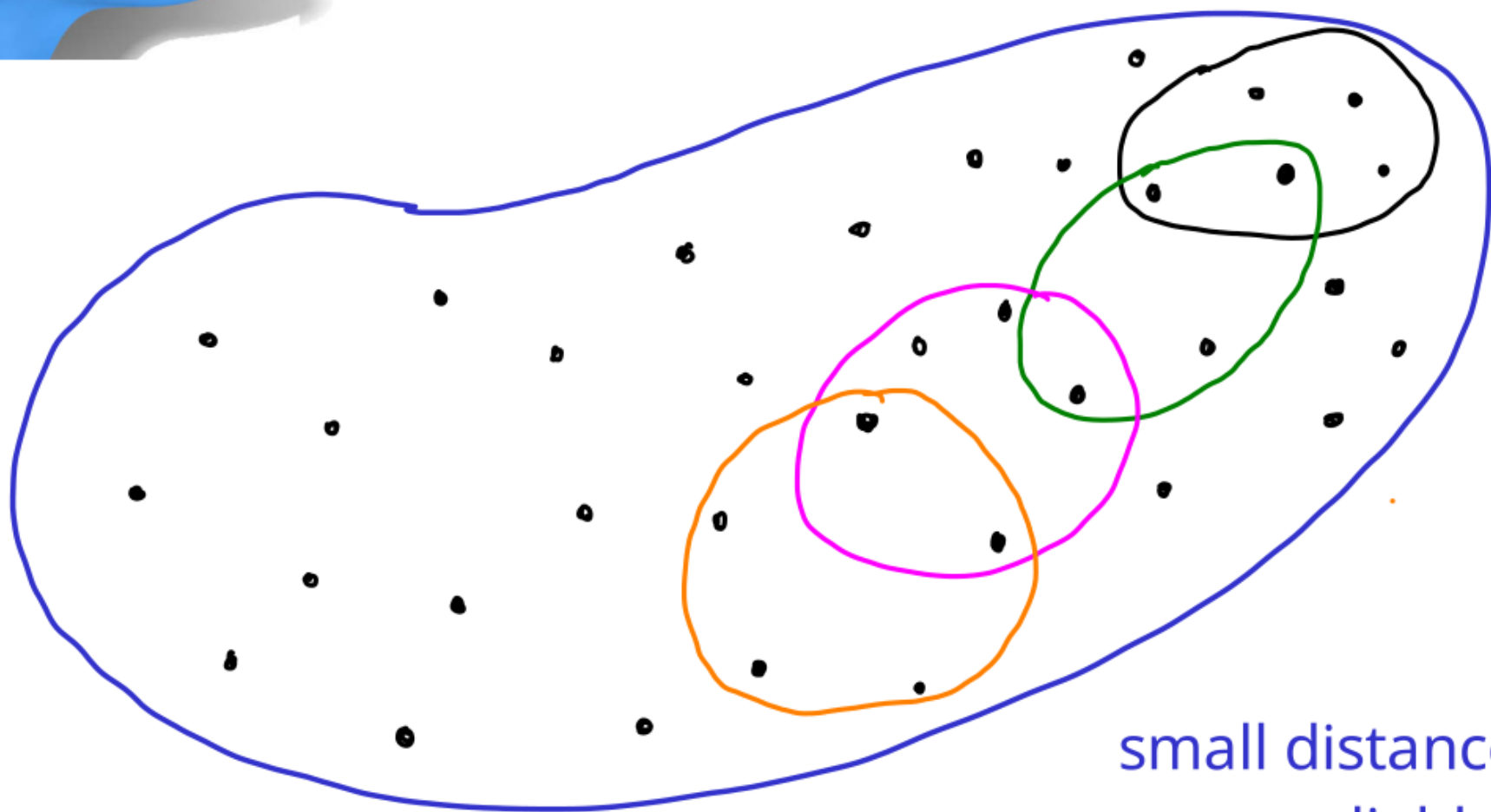
Data set



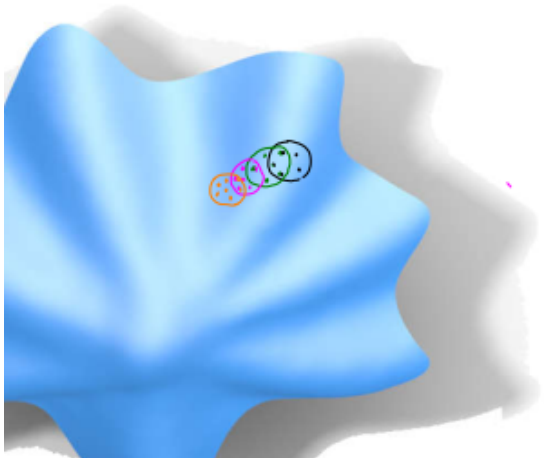
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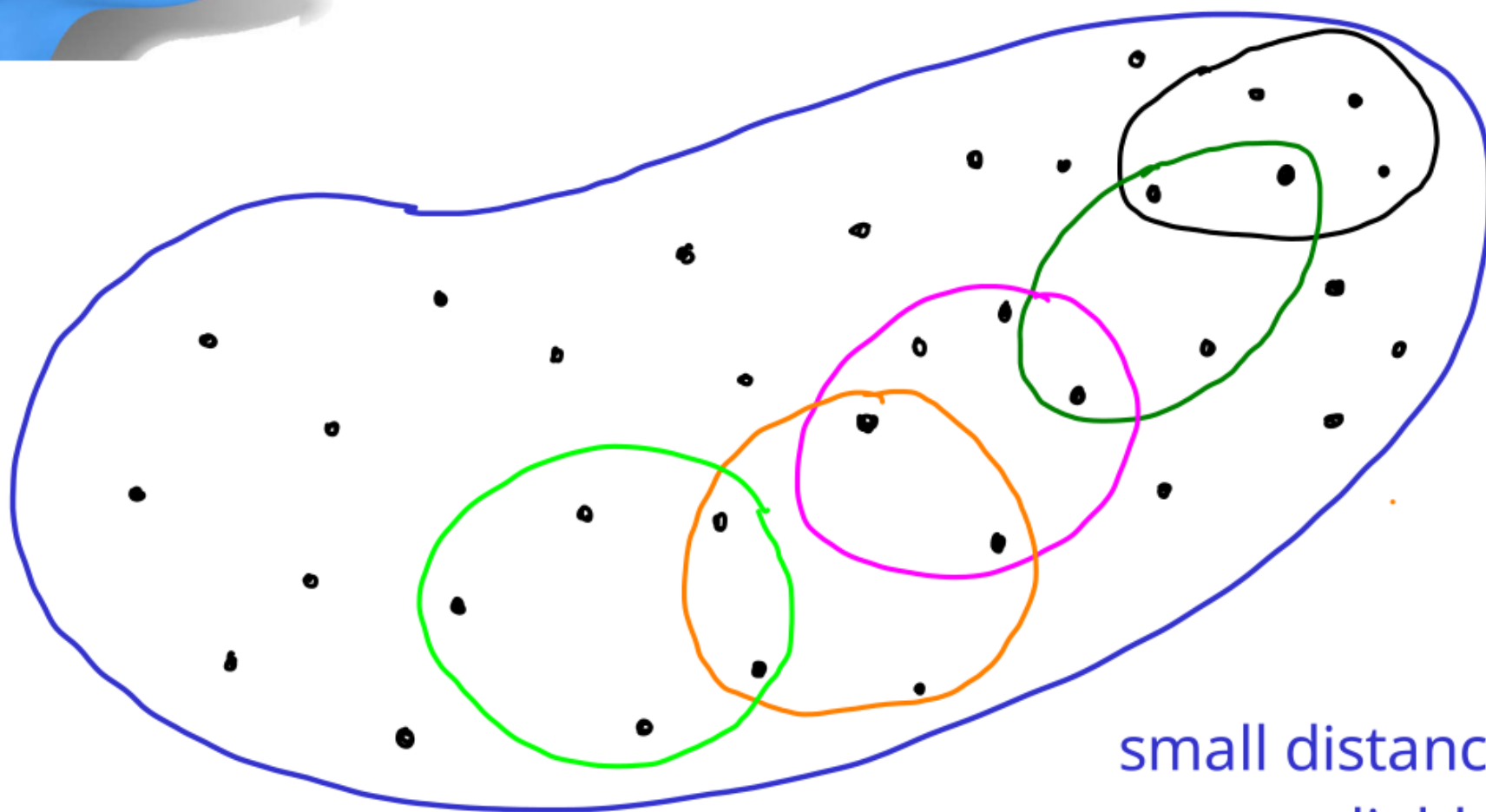
Data set



small distances are
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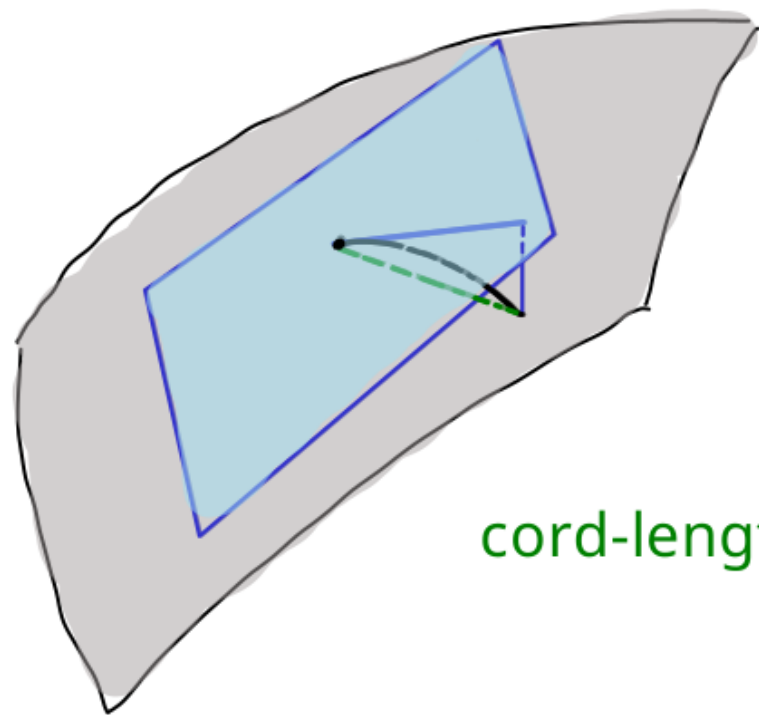
Data set



small distances are
more reliable!

How do this computationally?

build diffusion operator (matrix)



in first approximation, and
very locally,

cord-length $\simeq$ dist. on manifold

$\simeq$ proj. dist. on tangent plane

$\Rightarrow$ use standard expression for heat diffusion kernel

use standard expression for heat kernel

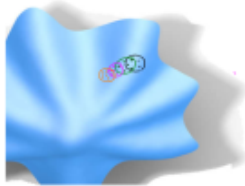
data points $P_i, P_j, \dots$

"dissimilarity" distance $\text{dist}(P_i, P_j) = d_{i,j}$

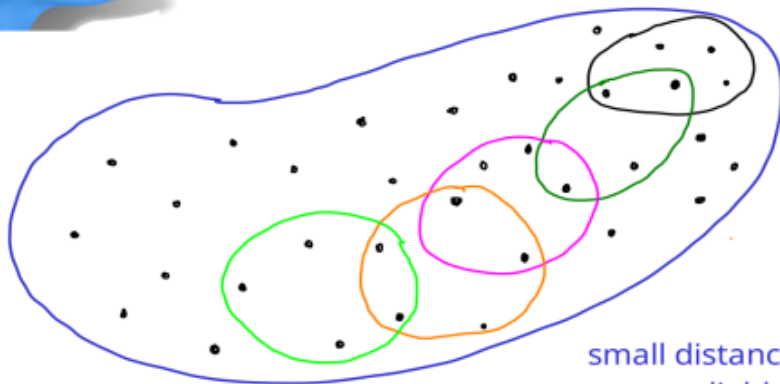
diffusion kernel in Euclidean space:

$$D_{t; i, j} = N_t e^{-d_{ij}^2 / 2t}$$

size of t measures extent of the diffusion



Data set



small distances are more reliable!

use standard expression for heat kernel

$$D_{t;i,j} = N_t e^{-d_{ij}^2 / 2t}$$

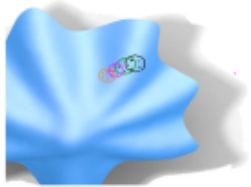


size of t indicates extent of the diffusion

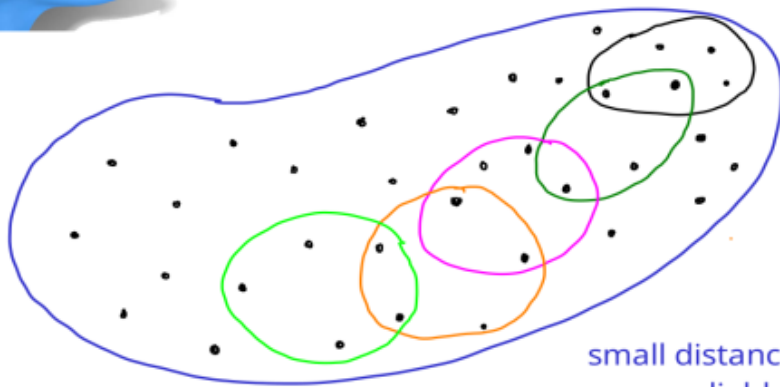
t needs to be sufficiently small

To "diffuse" further: consider powers of the diffusion matrix!

Normalization?



Data set

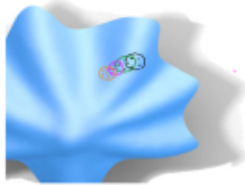


small distances are more reliable!

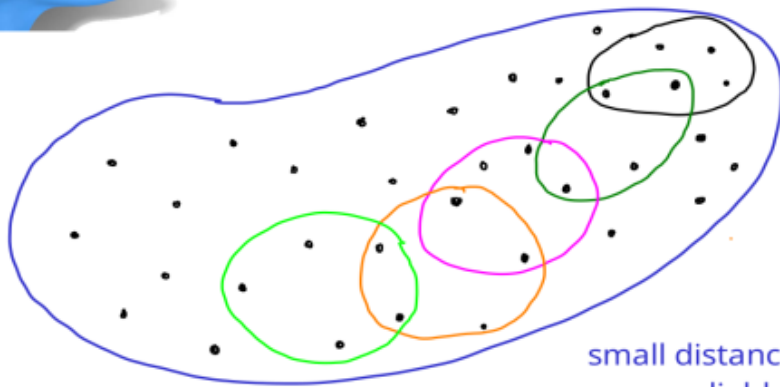
use standard expression for heat kernel

$$\mathbb{D}_{t;i,j} = N_t e^{-d_{ij}^2/2t}$$

Normalization?



Data set

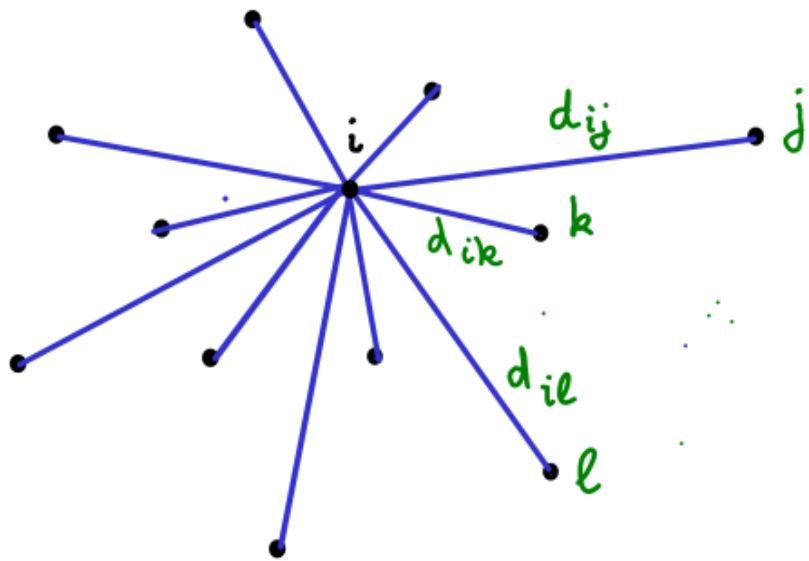


small distances are
more reliable!

Heat diffusion on a manifold
defines a random walk.

The approximation built from the
data should do likewise:

$$\sum_j \mathbb{D}_{t;i,j} = 1.$$
$$\Rightarrow N_t = \left(\sum_j e^{-d_{ij}^2/2t} \right)^{-1}$$



The data points and their distances define a weighted graph

$$e^{-d_{ij}^2/2t} = \mathcal{D}_{t;ij}$$

$$\tilde{\mathcal{D}}_{t;i,i} = \sum_j \mathcal{D}_{t;ij}$$

$$W_t = \tilde{\mathcal{D}}_t^{-1} \mathcal{D}_t$$

: defines a random walk on this graph

$\Rightarrow$ candidate for a discrete approximation of the diffusion on the manifold

How should one pick t ?

Not too large , not too small

$$e^{-d_{ij}^2/2t} = \mathcal{D}_{t;ij}$$

$$\tilde{\mathcal{D}}_{t;i,i} = \sum_j \mathcal{D}_{t;i,j}$$

$$W_t = \tilde{\mathcal{D}}_t^{-1} \mathcal{D}_t$$

candidate for a discrete
approximation of the diffusion
on the manifold

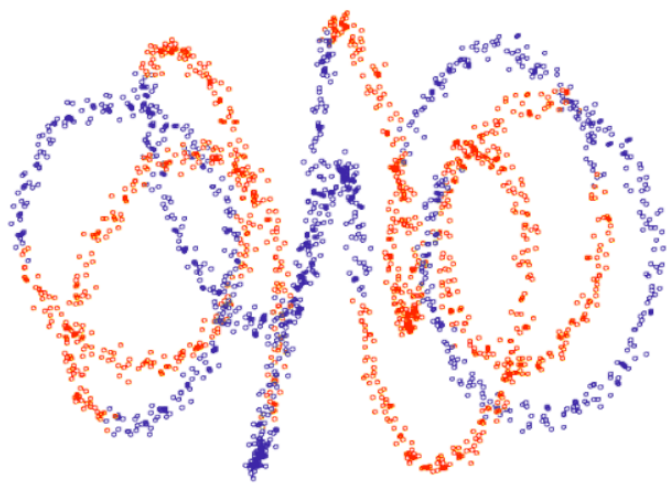
How should one pick t ?

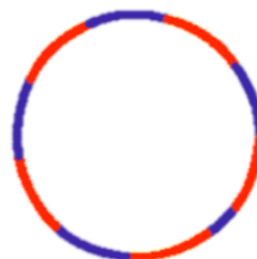
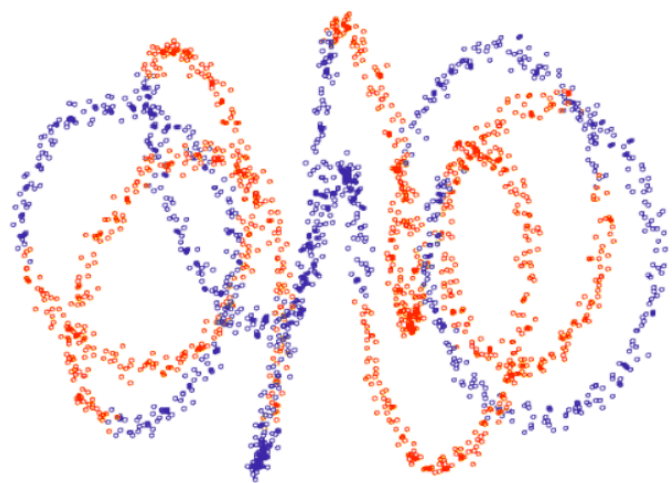
Not too large , not too small

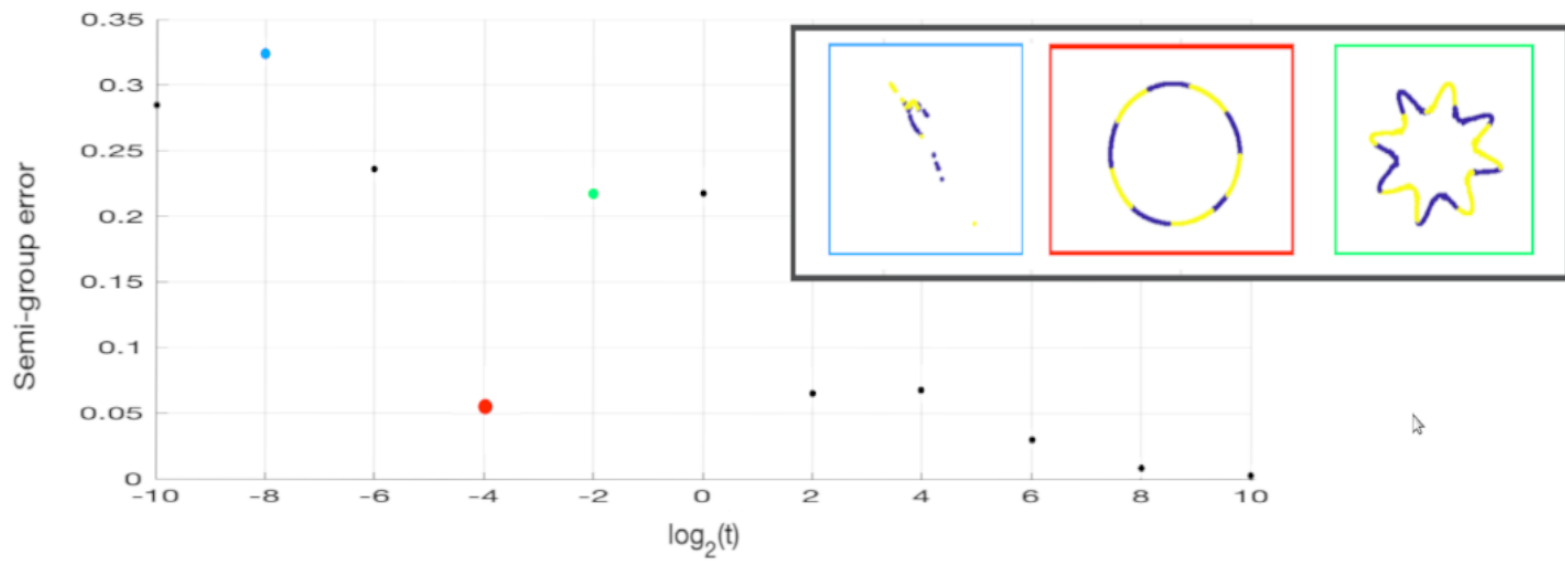
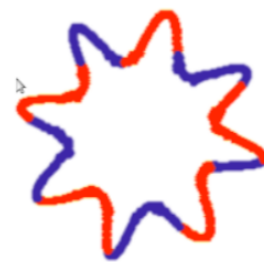
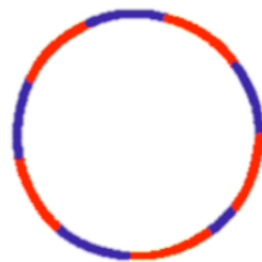
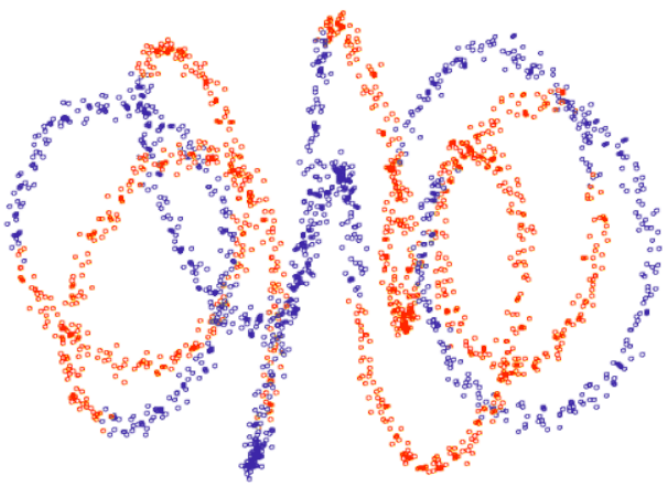
$W_t W_s$ should be close to W_{t+s}

$\| (W_t)^2 - W_{2t} \| \rightarrow$ should be small

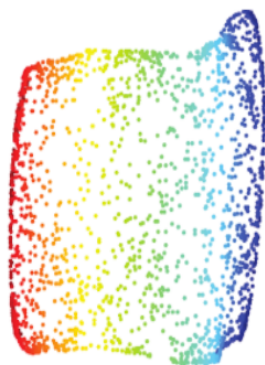
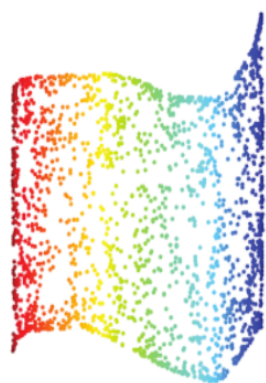
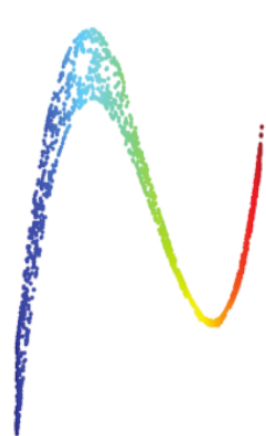
Shan Shan & ID





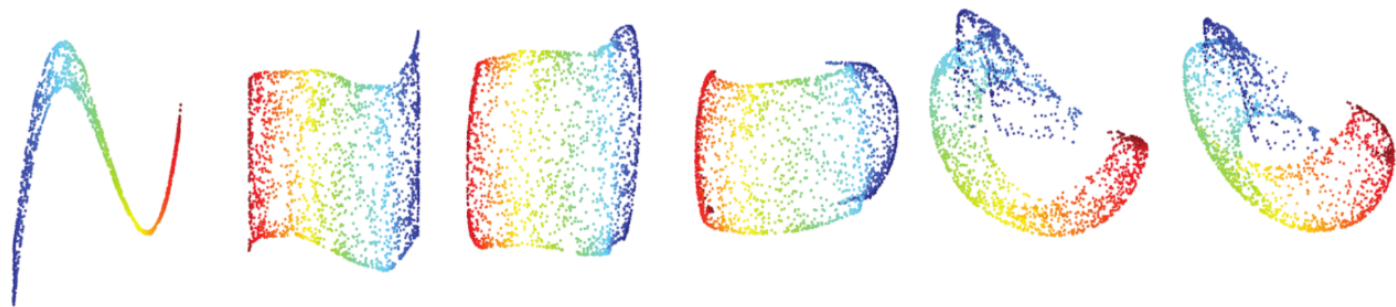




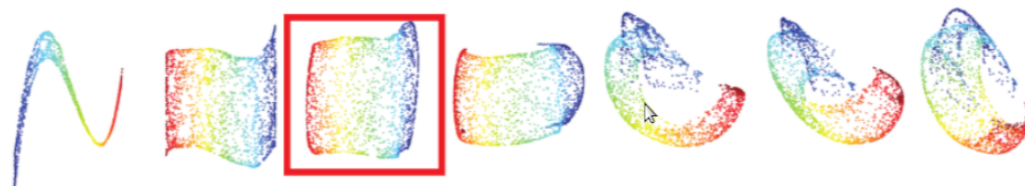




SGE



t



$$e^{-d_{ij}^2/2t} = \mathcal{D}_{t;ij}$$

$$\tilde{\mathcal{D}}_{t;i,i} = \sum_j \mathcal{D}_{t;ij}$$

$$W_t = \tilde{\mathcal{D}}_t^{-1} \mathcal{D}_t$$

candidate for a discrete
approximation of the diffusion
on the manifold

Next: spectral decomposition:

$$W_{t;ij} \cong \sum_l \omega_t \psi_{l;i} \psi_{l;j}$$

diffusion over time T: $\cong \sum_l (\omega_t)^{T/t} \psi_{l;i} \psi_{l;j}$

new parametrization for data points: $(\omega_t^\lambda \psi_{l;i})_{l=1}^L$

$$e^{-d_{ij}^2/2t} = \mathcal{D}_{t;ij}$$

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new parametrization for data points:

$$\left(\omega_t^\lambda \psi_{l,i} \right)_{l=1}^L$$

$\Rightarrow$ "diffusion distance"

$$d_{diff}^2(i,j) = \sum_{l=1}^L \omega_t^{2\lambda} |\psi_{l,i} - \psi_{l,j}|^2$$

.

So far: transition from a "dissimilarity distance" to a distance more governed by the entirety of the dataset.

ref: Belkin & Nyogi
Coifman, Lafon, Maggioni
Shan Shan

In this talk: application to data set of biological shapes
additional use of diffusion methods to provide even
better data-adapted distances.

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



Doug Boyer



Jukka Jernvall

More Precisely: biological morphologists



Study Teeth & Bones of
extant & extinct animals

still live today

fossils

Collaborators



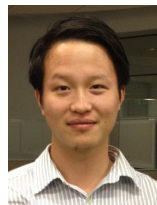
Rima Alaifari
ETH Zürich



Doug Boyer
Duke



Ingrid Daubechies
Duke



Tingran Gao
Duke



Yaron Lipman
Weizmann



Roi Poranne
ETH Zürich



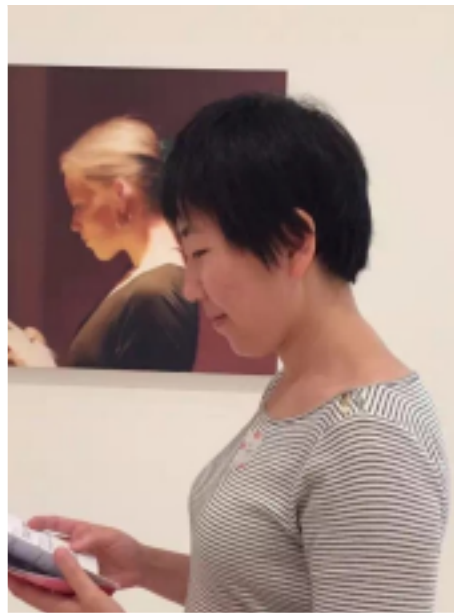
Jesús Puente
J.P. Morgan



Robert Ravier
Duke



Shahar Kovalsky



Shan Shan



Nadav Dym



Chen-Yun Lin



Shira Faigenbaum



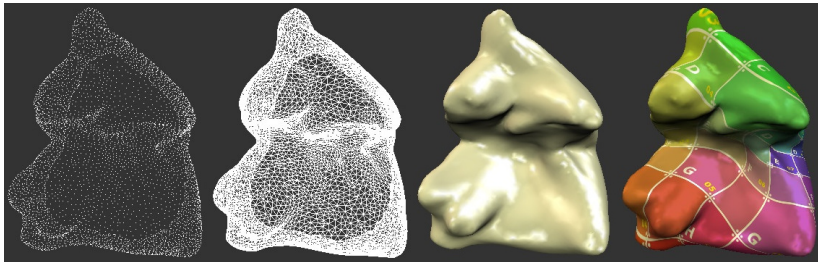
Alex Winn

First: project on “complexity” of teeth

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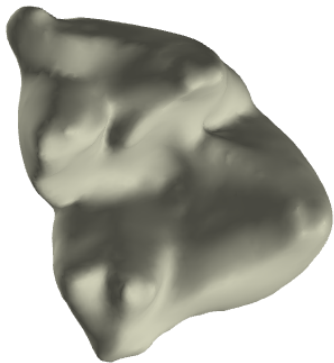
Then: find automatic way to compute Procrustes distances
between surfaces — without landmarks

Data Acquisition



Surface reconstructed from μ CT-scanned voxel data

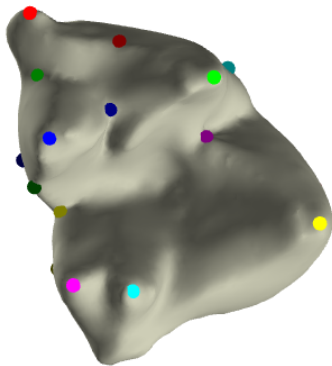
Geometric Morphometrics



second mandibular molar of a Philippine flying lemur

- Manually put k landmarks

Geometric Morphometrics

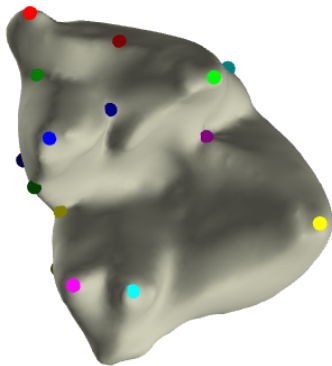


second mandibular molar of a Philippine flying lemur

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$$p_1, p_2, \dots, p_k$$

Geometric Morphometrics



second mandibular molar of a Philippine flying lemur

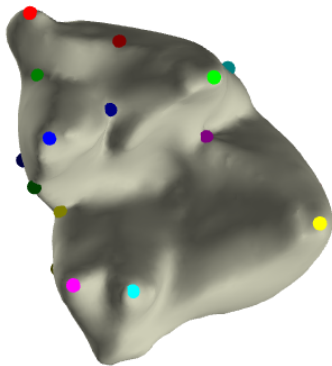
- Manually put k **landmarks**

$$p_1, p_2, \dots, p_k$$

- Use **spatial** coordinates of the landmarks as features

$$p_j = (x_j, y_j, z_j), j = 1, \dots, k$$

Geometric Morphometrics



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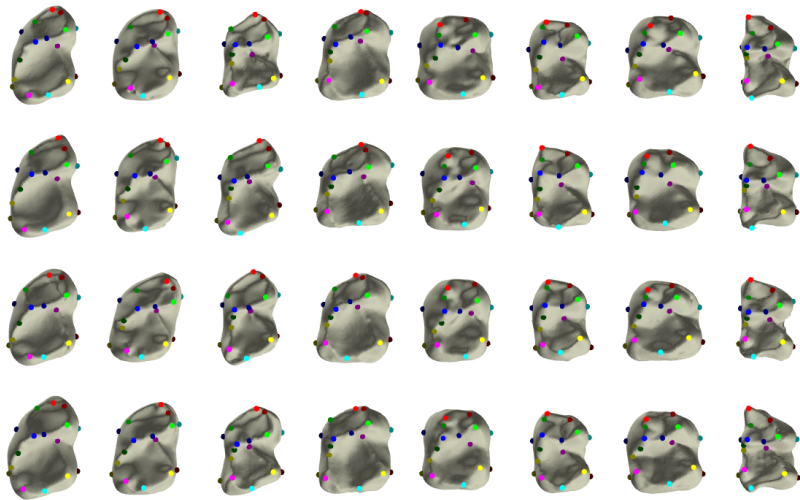
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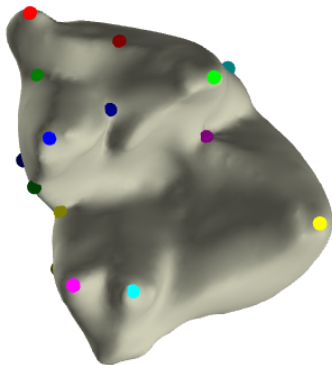
$$p_j = (x_j, y_j, z_j), \quad j = 1, \dots, k$$

- Represent a shape in $\mathbb{R}^{3 \times k}$

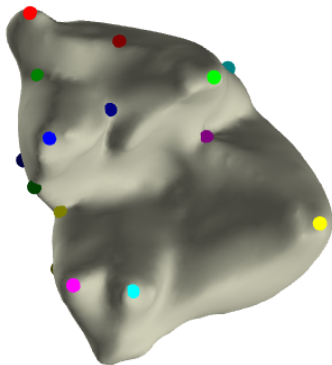
The *Shape Space* of k landmarks in $\mathbb{R}^3$



Geometric Morphometrics: Limitation of Landmarks

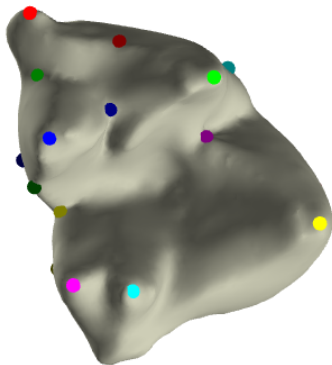


Geometric Morphometrics: Limitation of Landmarks



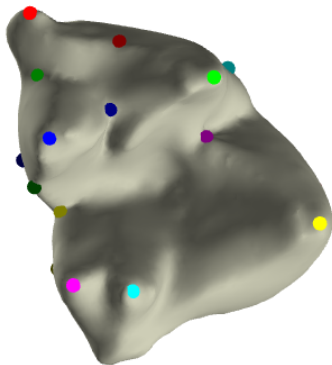
- **Landmark Placement:** tedious and time-consuming

Geometric Morphometrics: Limitation of Landmarks



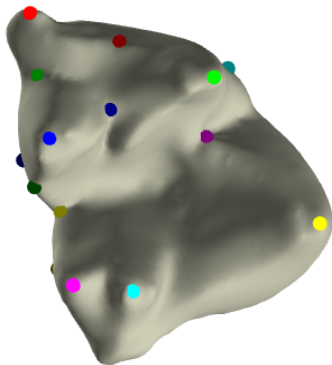
- **Landmark Placement:** tedious and time-consuming
- **Fixed Number of Landmarks:** lack of flexibility

Geometric Morphometrics: Limitation of Landmarks



- **Landmark Placement:** tedious and time-consuming
- **Fixed Number of Landmarks:** lack of flexibility
- **Domain Knowledge:** high degree of expertise needed, not easily accessible

Geometric Morphometrics: Limitation of Landmarks



- **Landmark Placement:** tedious and time-consuming
- **Fixed Number of Landmarks:** lack of flexibility
- **Domain Knowledge:** high degree of expertise needed, not easily accessible
- **Subjectivity:** debates exist even among experts

First: project on “complexity” of teeth

Then: find automatic way to compute Procrustes distances
between surfaces — without landmarks



Landmarked Teeth $\longrightarrow$

$$d_{Procrustes}^2(S_1, S_2) = \min_{R \text{ rigid tr.}} \sum_{j=1}^J \|R(x_j) - y_j\|^2$$

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



First: project on “complexity” of teeth

Then: find automatic way to compute Procrustes distances between surfaces — without landmarks

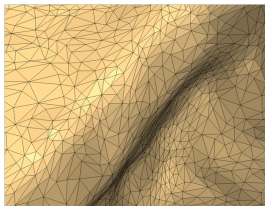
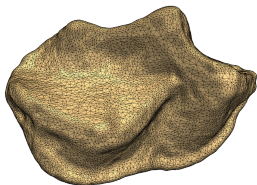


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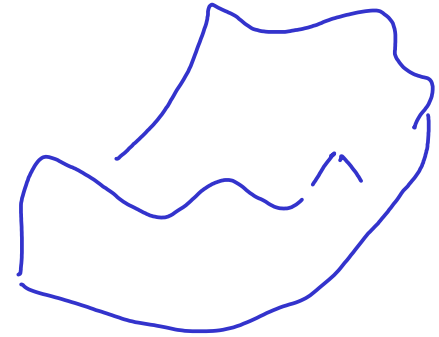
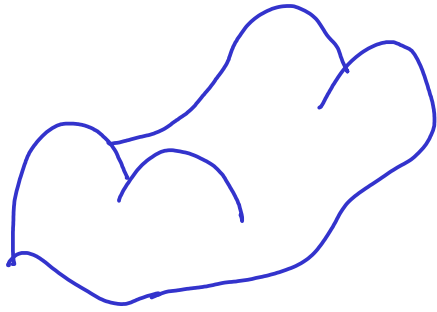
Find way to compute a distance that does as well, for biological purposes, as Procrustes distance, based on expert-placed landmarks, **automatically**?

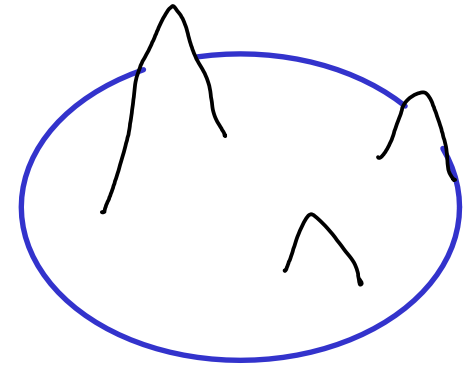
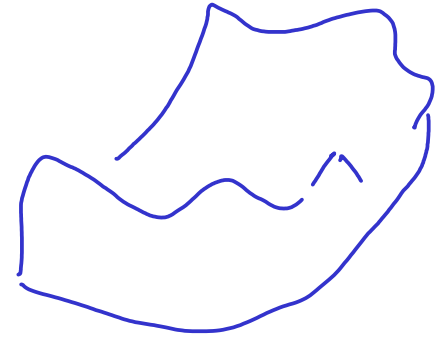
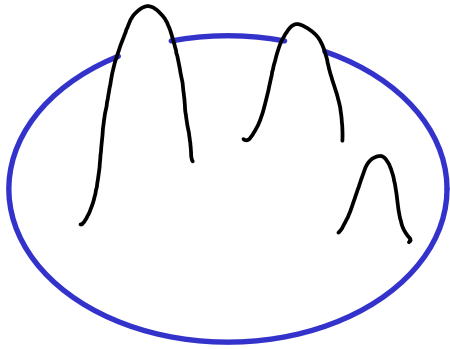
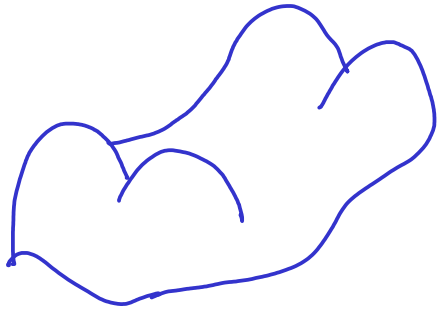
examples: finely discretized triangulated surfaces

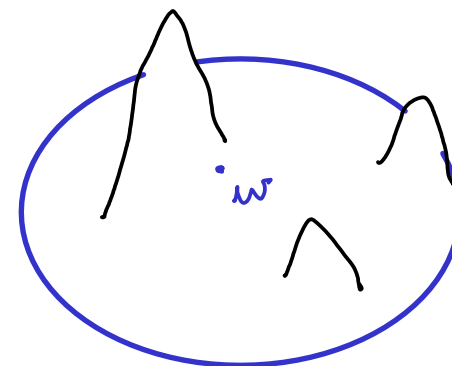
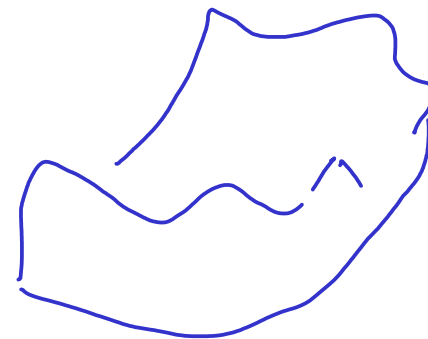
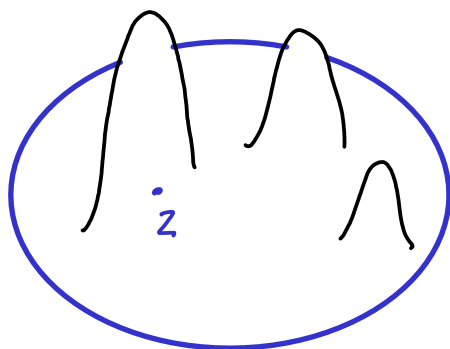
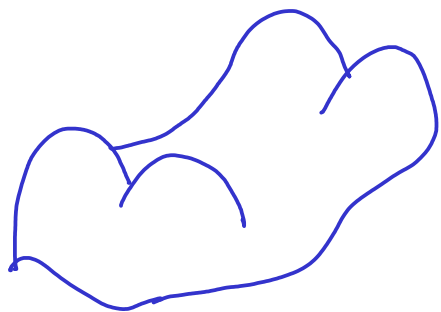


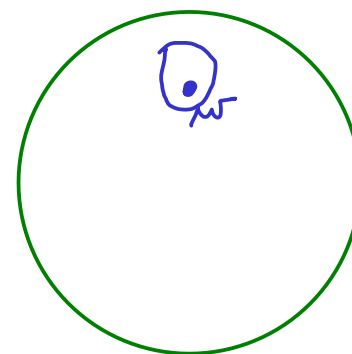
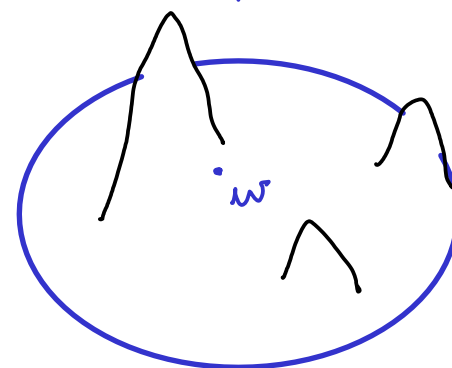
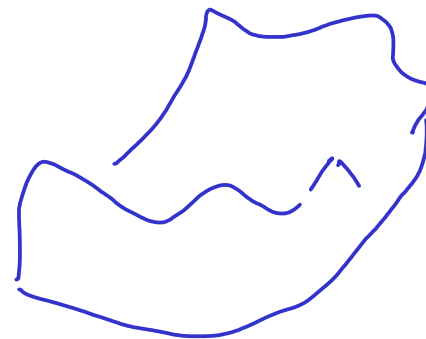
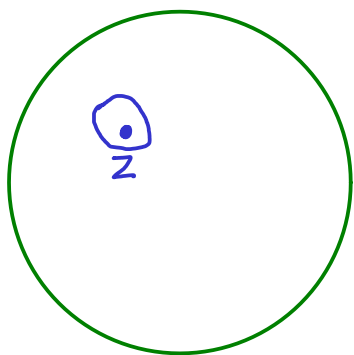
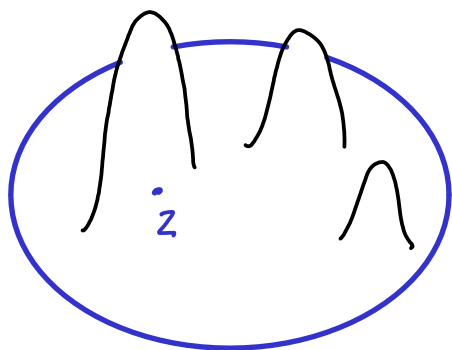
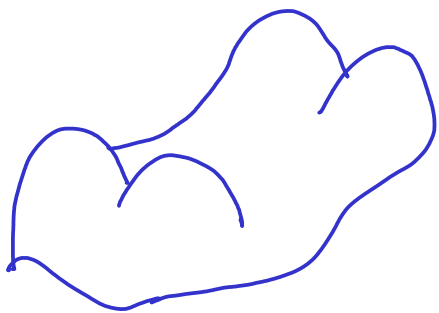
We defined 2 different distances

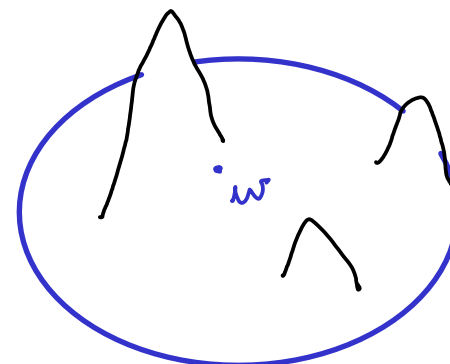
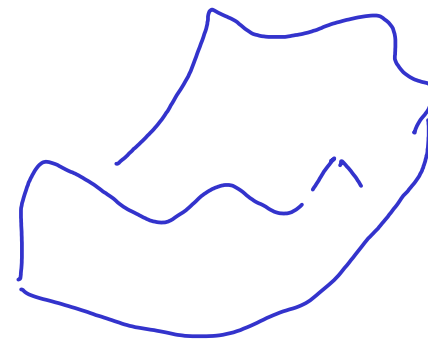
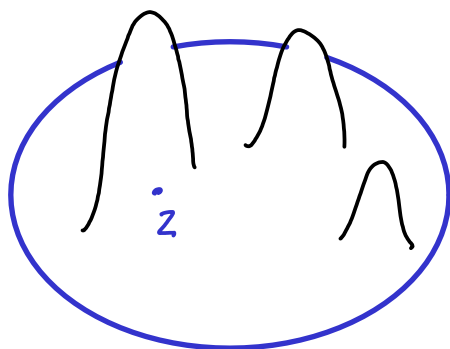
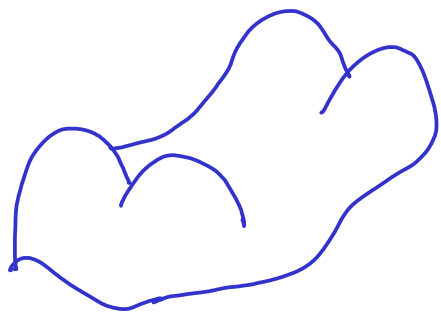
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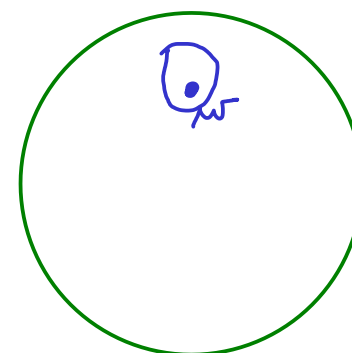
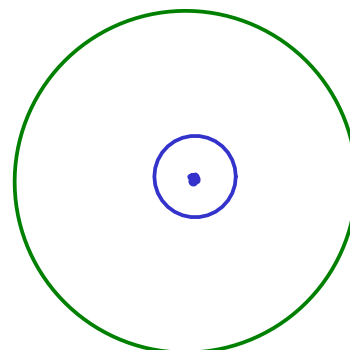
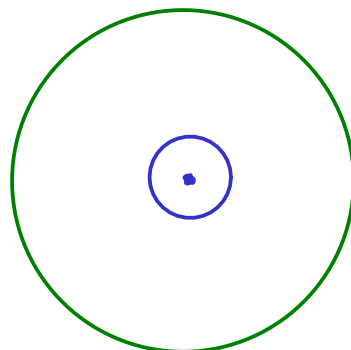
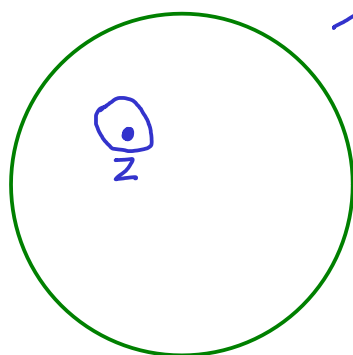


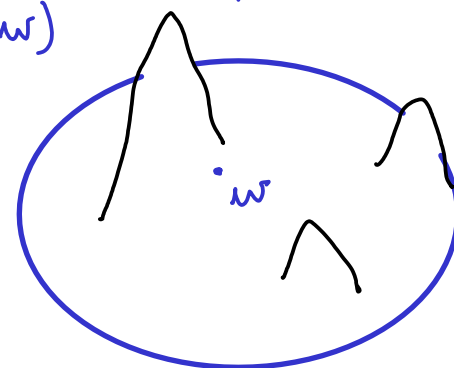
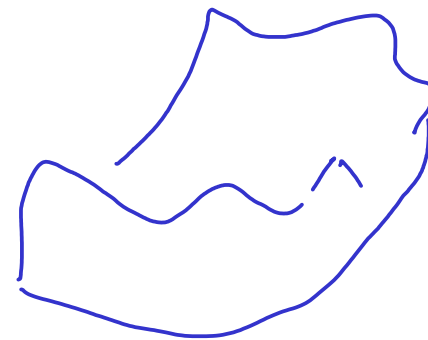
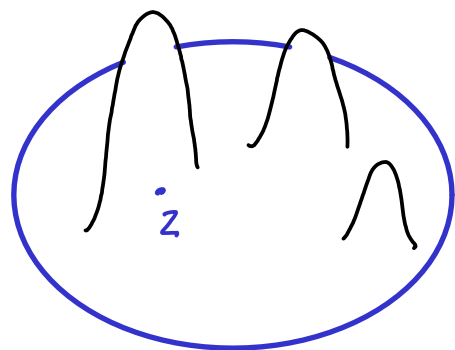
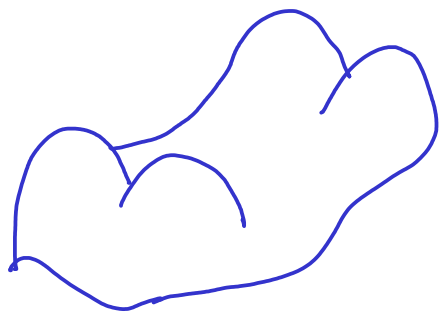






$$d_{R}^{\mu, \nu}(z, w)$$

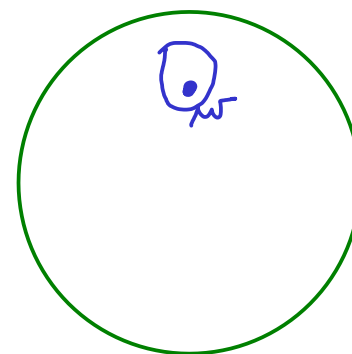
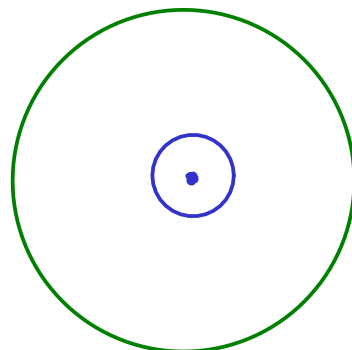
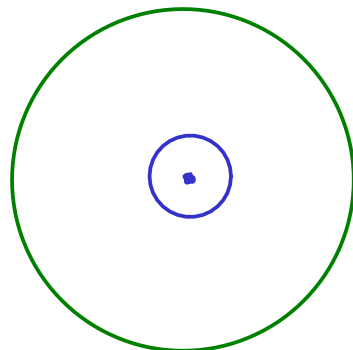
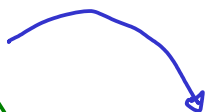
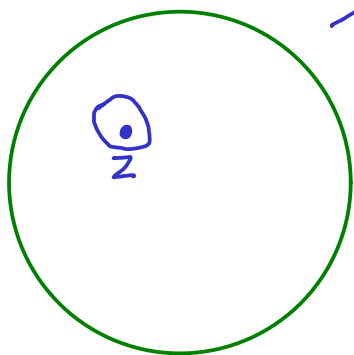




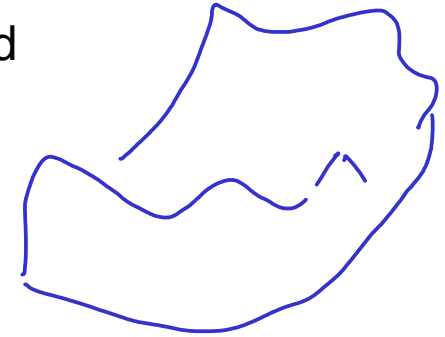
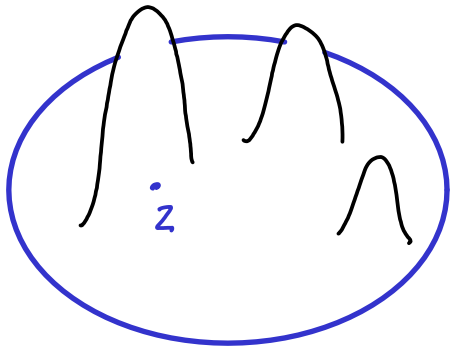
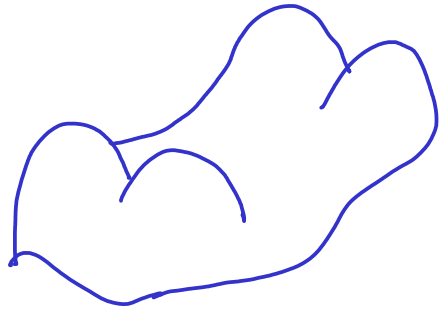
$$\mathbb{D}(S_1, S_2) = \inf_{\pi \in \Pi(\mu, \nu)} \int d_R^{\mu, \nu}(z, w) d\pi(z, w)$$



$$d_R^{\mu, \nu}(z, w)$$



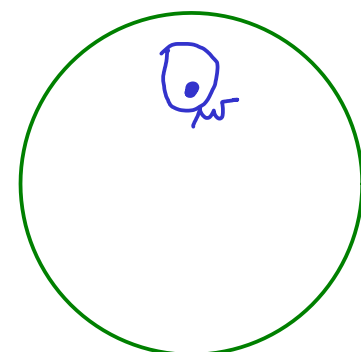
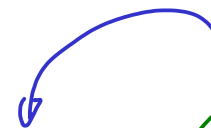
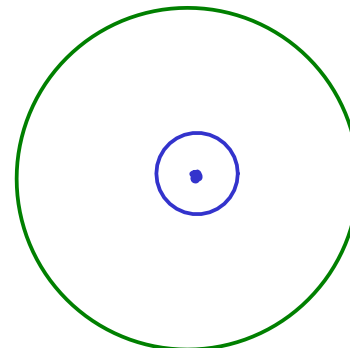
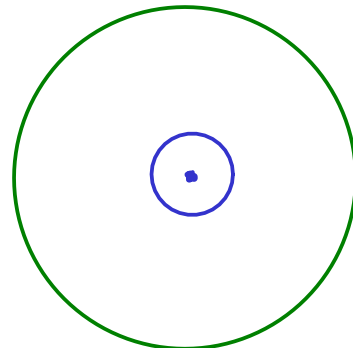
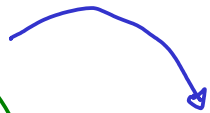
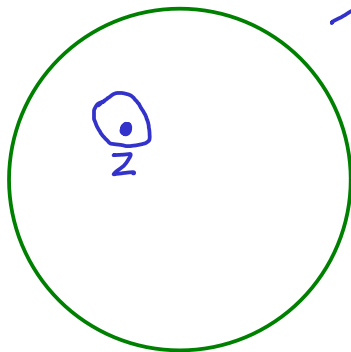
conformal Wasserstein neighborhood
distance



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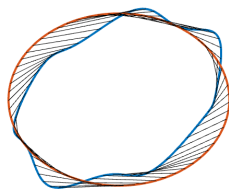
$$d_R^{\mu, \nu}(z, w)$$



Continuous Procrustes Distance (cPD)

$$D_{\text{cP}}(S_1, S_2) = \left(\int_{S_1} \|x - \mathcal{C}(x)\|^2 d\text{vol}_{S_1}(x) \right)^{\frac{1}{2}},$$

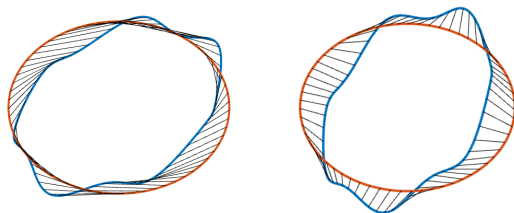
where $\mathcal{C} : S_1 \rightarrow S_2$ is an area-preserving diffeomorphism.



Continuous Procrustes Distance (cPD)

$$D_{\text{cP}}(S_1, S_2) = \left(\inf_{R \in \mathbb{E}(3)} \int_{S_1} \|R(x) - \mathcal{C}(x)\|^2 d\text{vol}_{S_1}(x) \right)^{\frac{1}{2}},$$

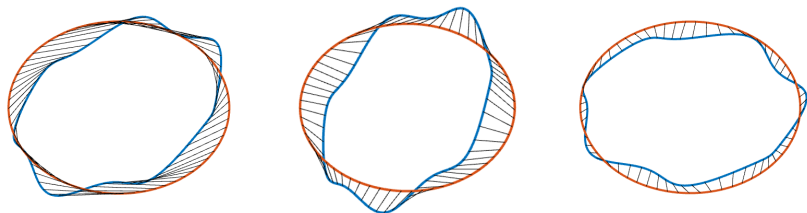
where $\mathcal{C} : S_1 \rightarrow S_2$ is an **area-preserving diffeomorphism**, and $\mathbb{E}_3$ is the Euclidean group on $\mathbb{R}^3$.



Continuous Procrustes Distance (cPD)

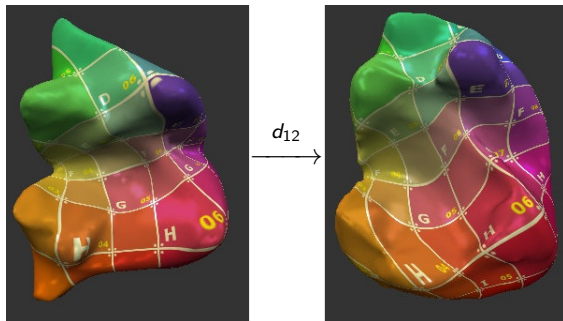
$$D_{\text{cP}}(S_1, S_2) = \left(\inf_{C \in \mathcal{A}(S_1, S_2)} \inf_{R \in \mathbb{E}(3)} \int_{S_1} \|R(x) - C(x)\|^2 d\text{vol}_{S_1}(x) \right)^{\frac{1}{2}},$$

where $\mathcal{A}(S_1, S_2)$ is the set of **area-preserving diffeomorphisms** between S_1 and S_2 , and $\mathbb{E}_3$ is the Euclidean group on $\mathbb{R}^3$.



Continuous Procrustes Distance (cPD)

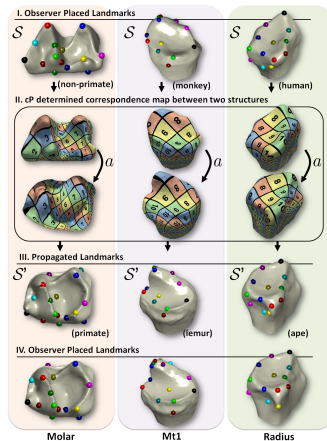
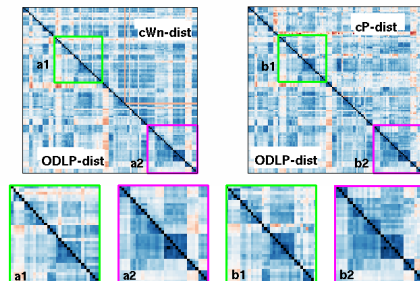
$$d_{cP}(S_1, S_2) = \inf_{C \in \mathcal{A}} \inf_{R \in \mathbb{E}_3} \left(\int_{S_1} \|R(x) - C(x)\|^2 d\text{vol}_{S_1}(x) \right)^{1/2}$$



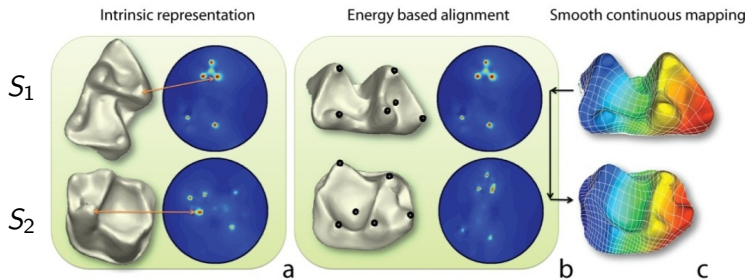
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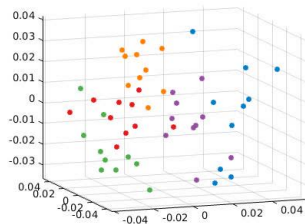
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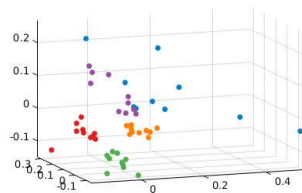
Bypass Explicit Feature Extraction



MDS for cPD & DD

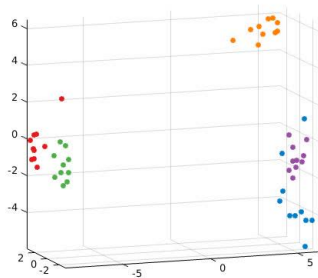


cPD

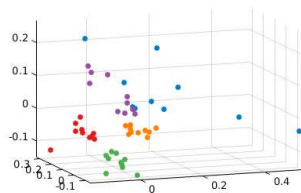


DD

Even better can be obtained!

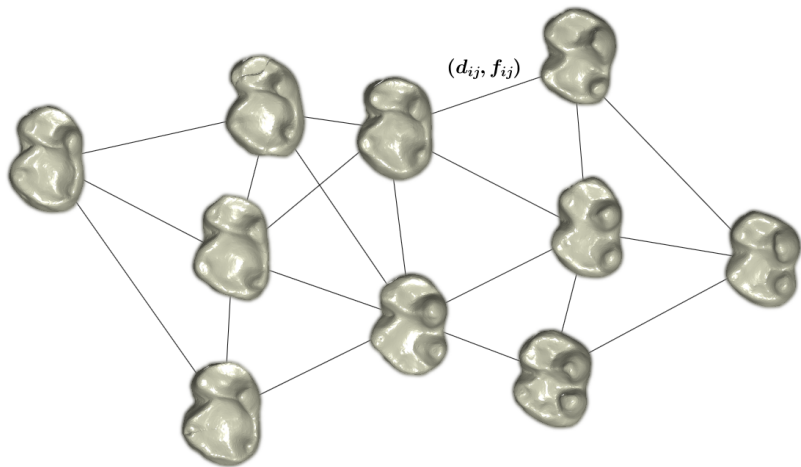


HBDD



DD

to get Diffusion Distance : used local distances
knitted together
→ spectral parametrization
→ distance.



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mappings were used only to obtain numerical
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but they can do much more for us!

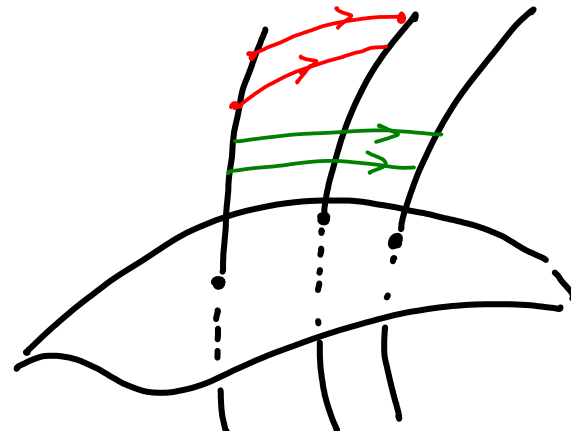
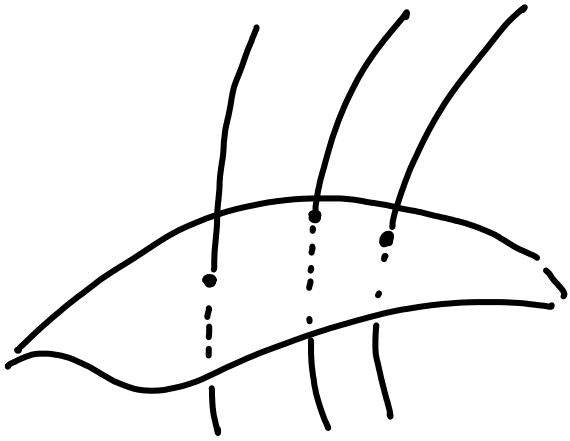
in fact: we have a fiber bundle.
(because of the mappings)

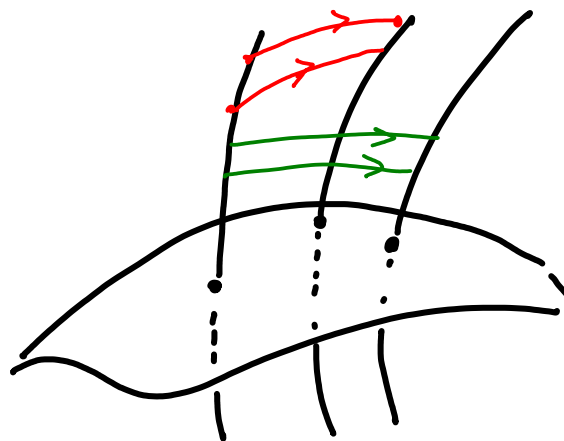
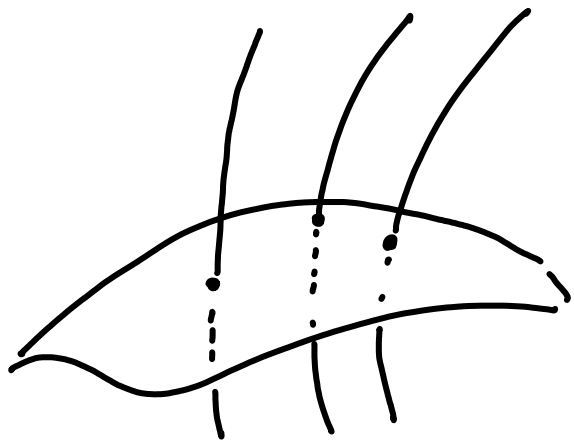
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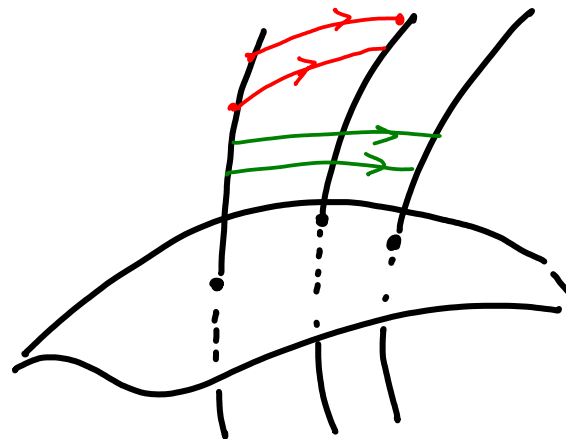
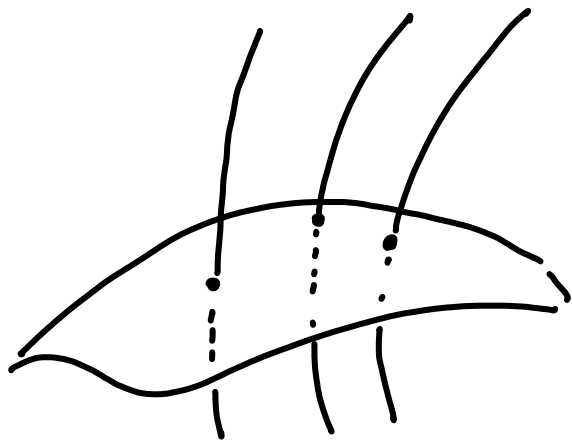




Connection.



family of mappings between fibers



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family of mappings between fibers

Tigran Gao: use these to define a much more detailed diffusion structure on the higher-dimensional object
 → "project" at a later stage to obtain "horizontal" part of diffusion.

Horizontal Random Walk on a Fibre Bundle

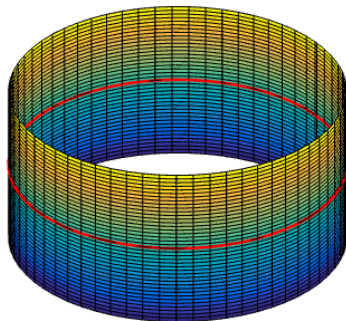
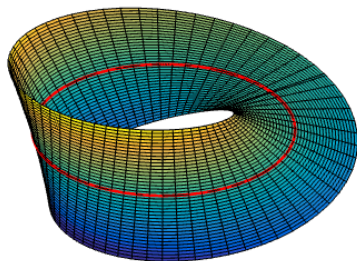
Fibre Bundle $\mathcal{E} = (E, M, F, \pi)$

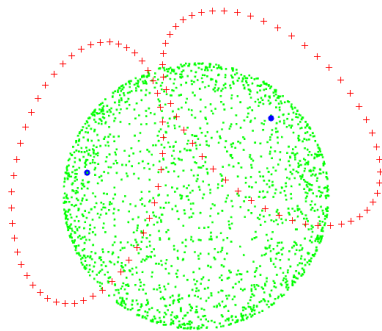
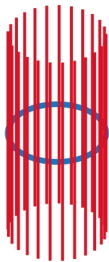
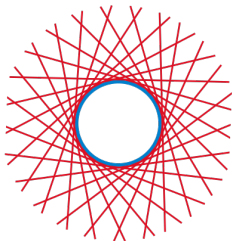
- ▶ E : *total* manifold
- ▶ M : *base* manifold
- ▶ $\pi : E \rightarrow M$: smooth surjective map (*bundle projection*)
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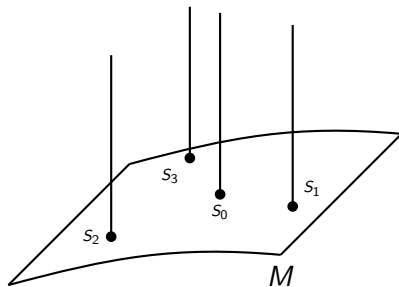
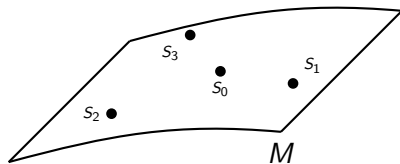




Horizontal Random Walk on a Fibre Bundle

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

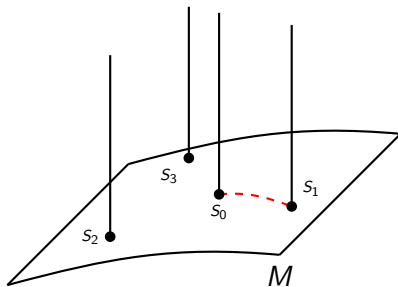
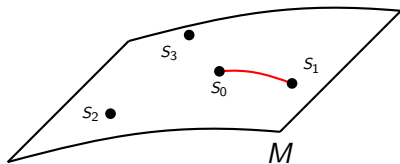
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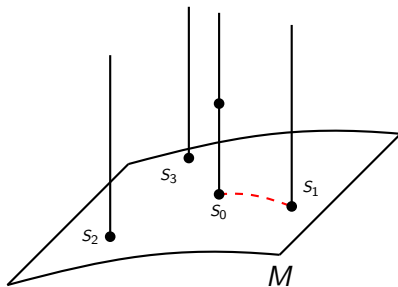
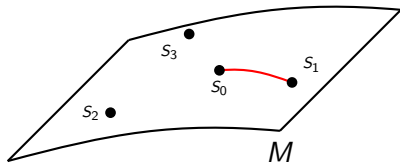
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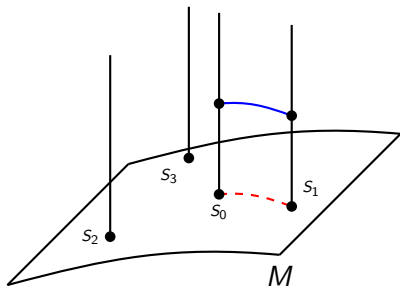
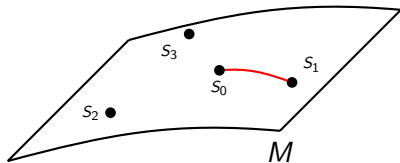
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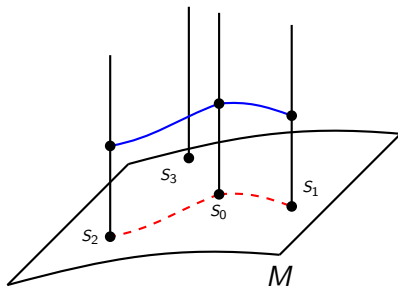
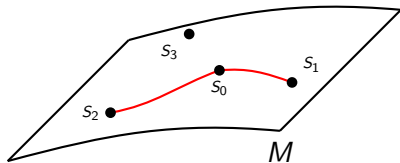
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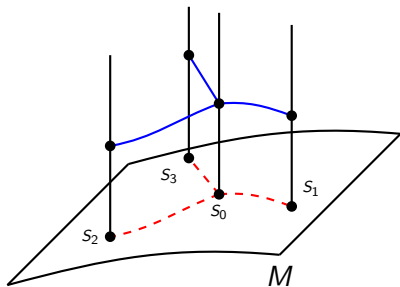
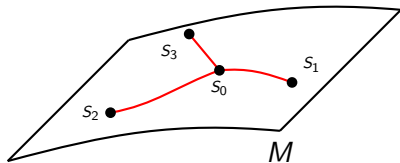
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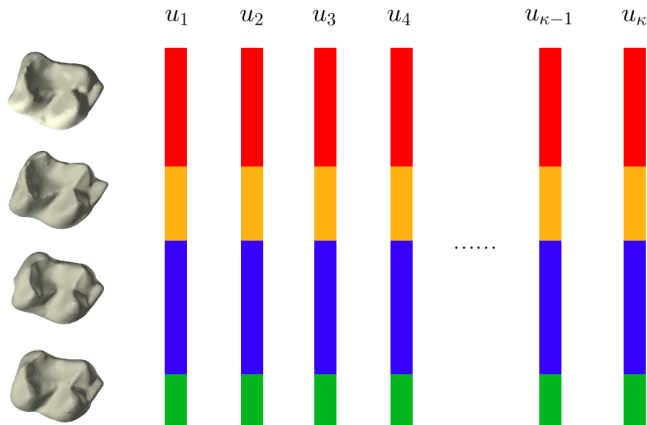
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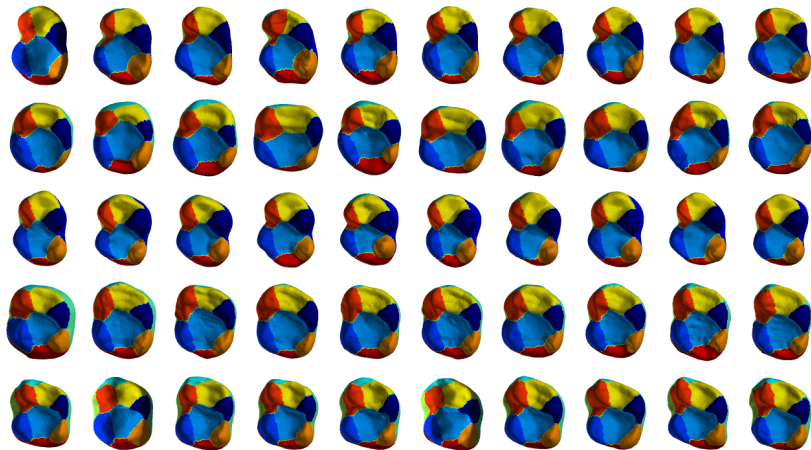
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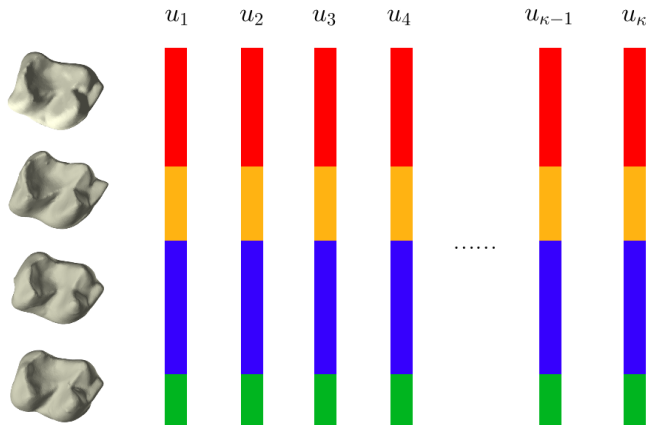
Horizontal Diffusion Maps: Embedding the Entire Bundle



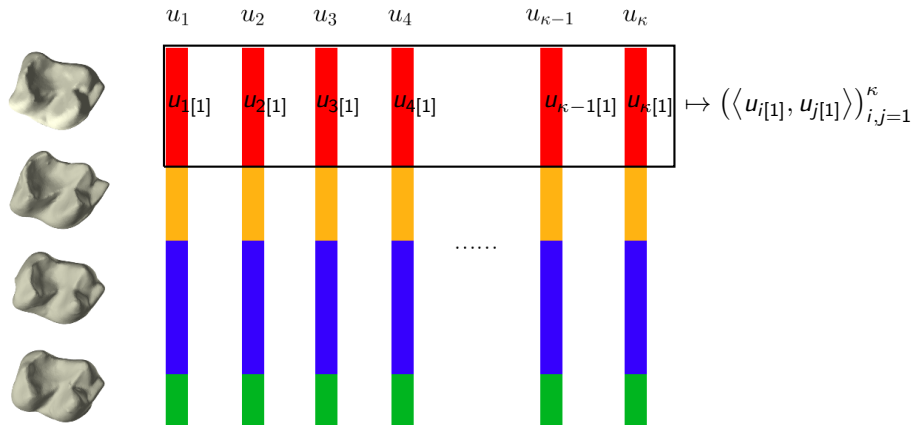
Automatic Landmarking — Interpretability



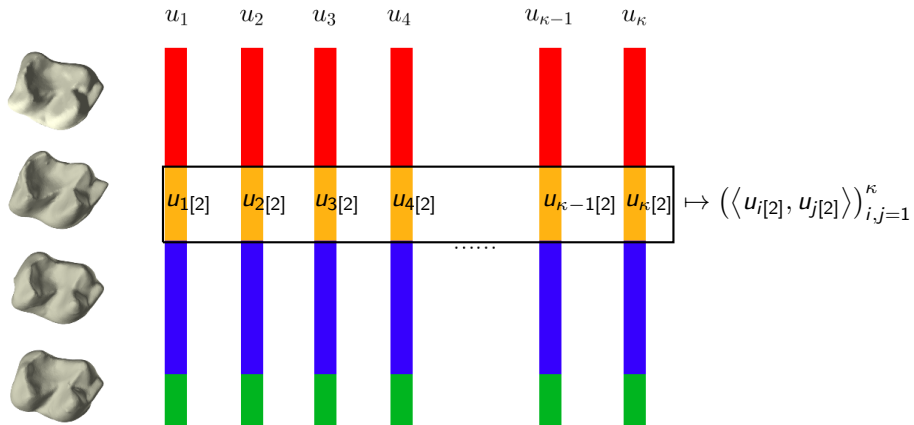
Horizontal Diffusion Maps: Embedding the Base Manifold



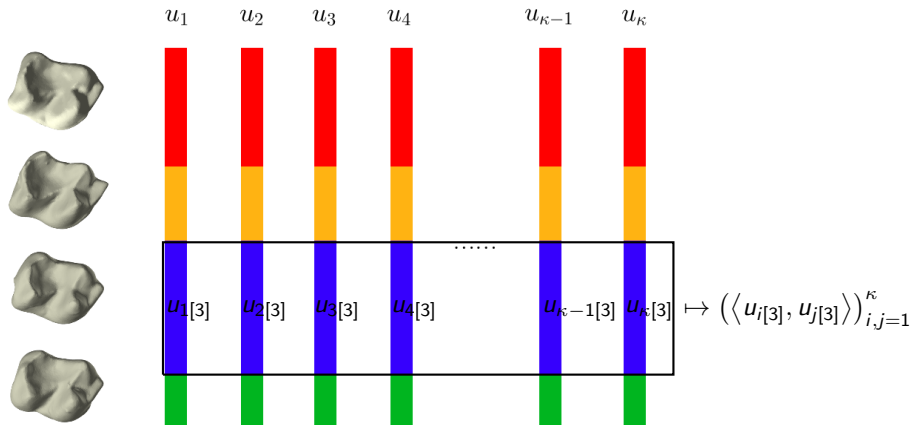
Horizontal Diffusion Maps: Embedding the Base Manifold



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Horizontal Diffusion Maps: Embedding the Base Manifold



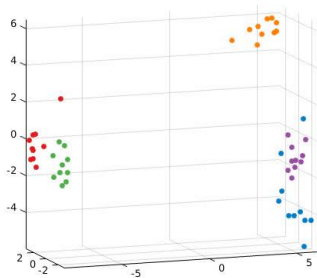
(j, p)
 $\swarrow$ point p on
 surface S_j

$$\rightarrow \left(u_k(j, p) \right)_{k=1}^K$$

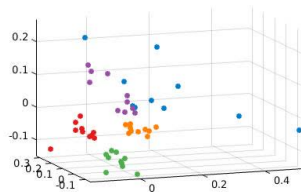
$$M_{j;k,\ell} = \sum_P w_{\ell,k}^\lambda u_k(j,P) \overline{u_\ell(j,P)}$$

$$\text{dist}(s_j, s_{j'}) = \|M_j - M_{j'}\|.$$

Species Clustering

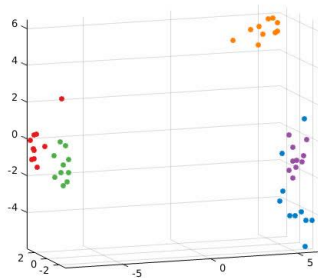


Horizontal Base Diffusion Distance (**with** Maps)

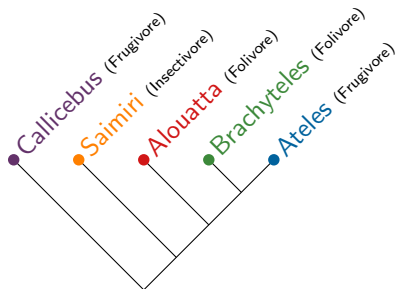


Diffusion Distance (**without** Maps)

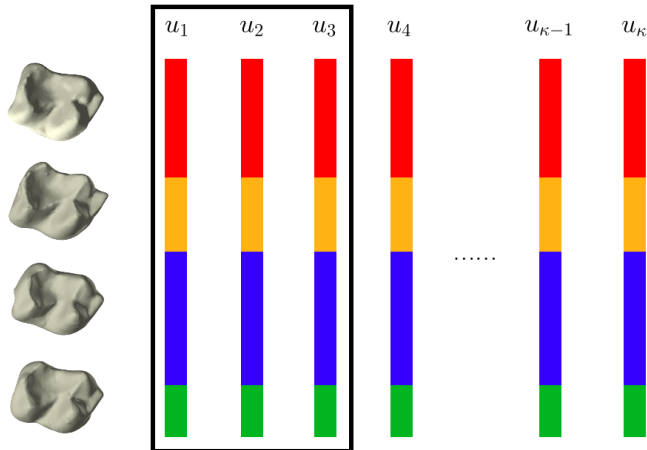
Species Clustering



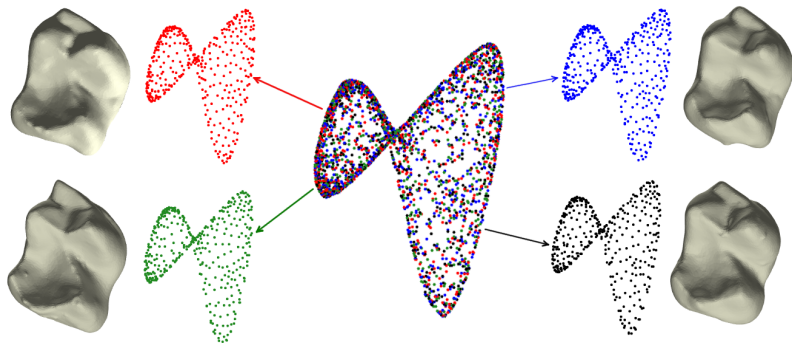
Horizontal Base Diffusion Distance (with Maps)

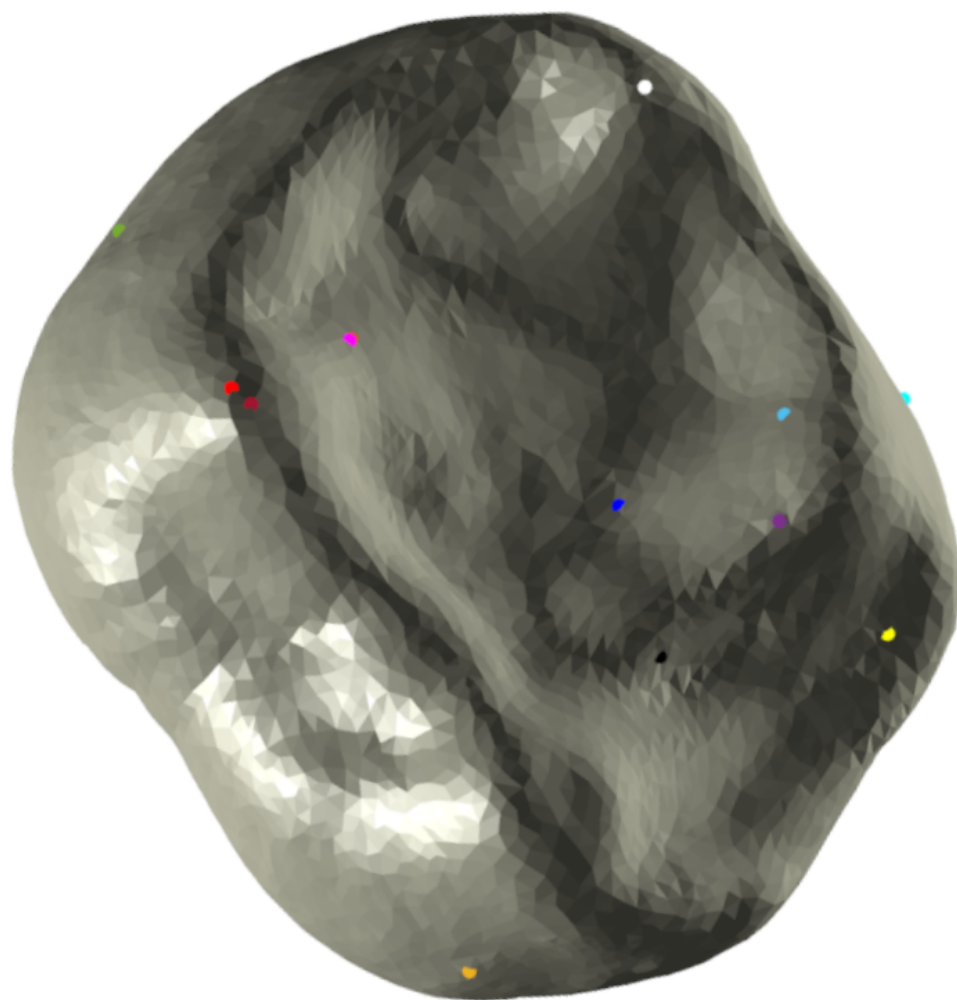
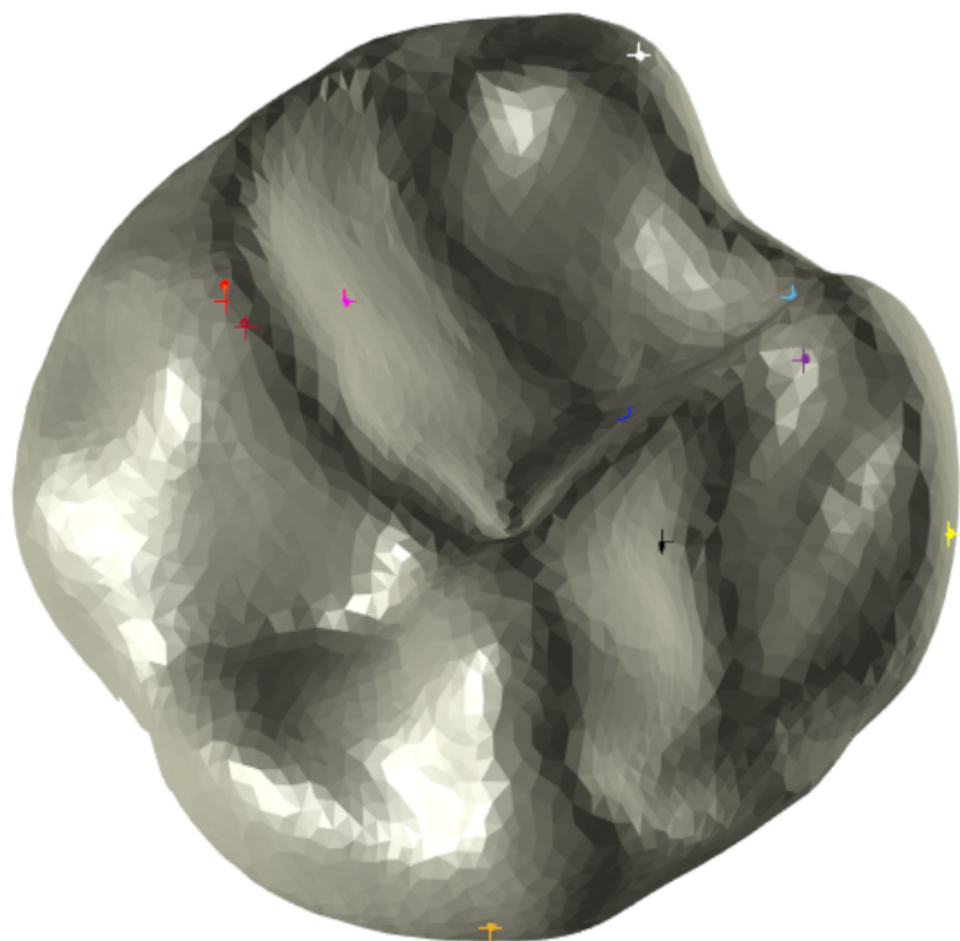


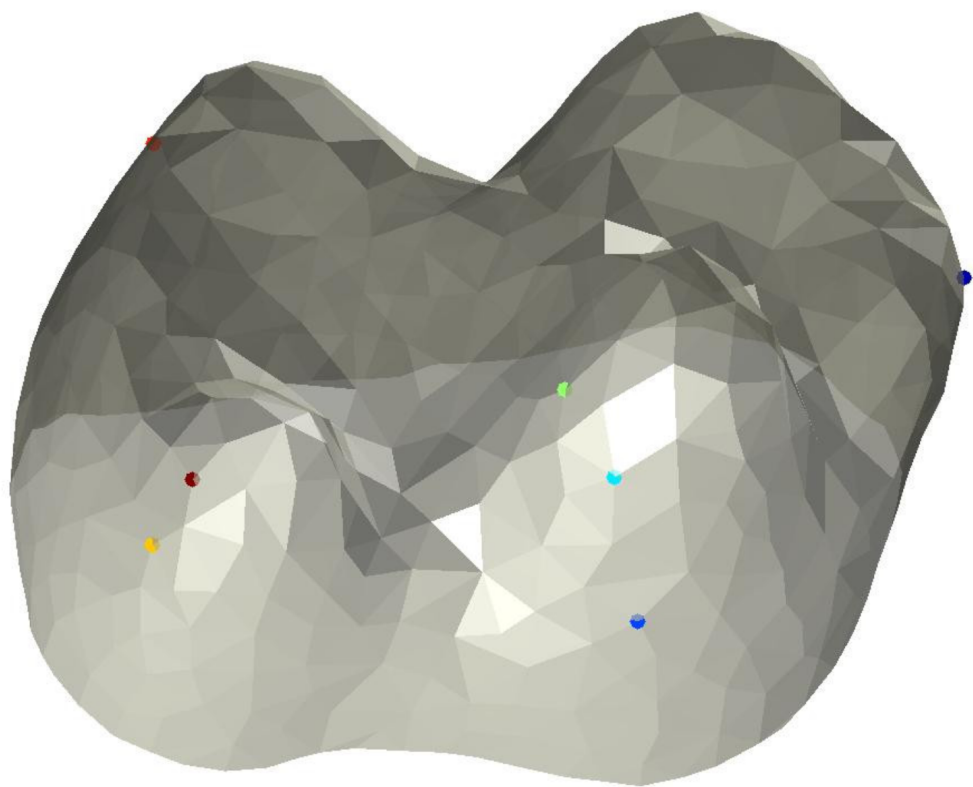
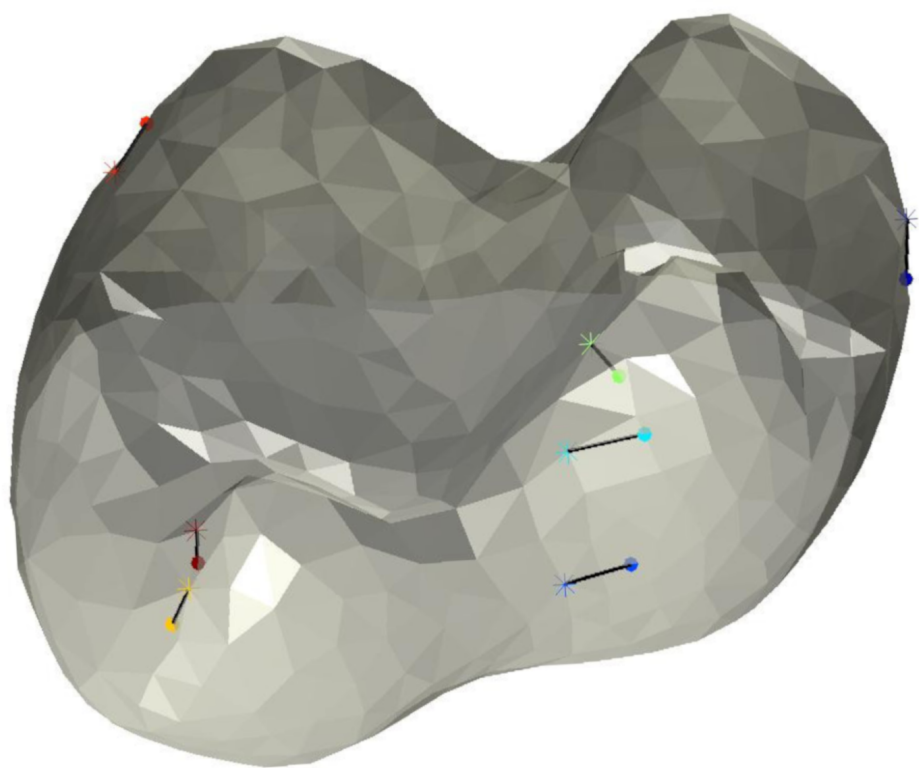
Horizontal Diffusion Maps: Embedding the Entire Bundle



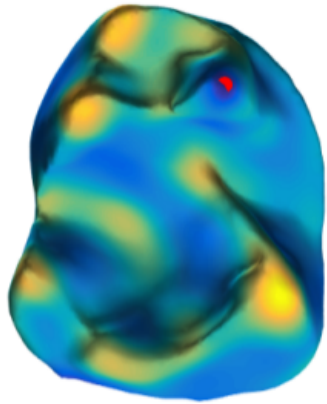
Horizontal Diffusion Maps



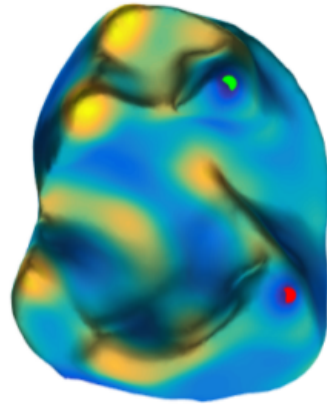




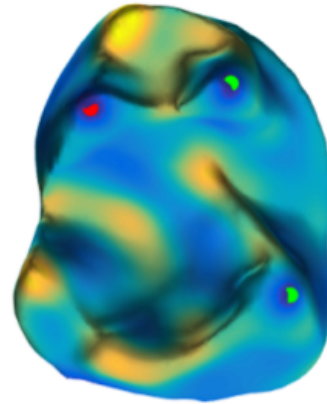
Step 1



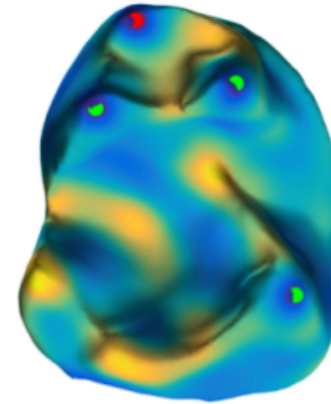
Step 2



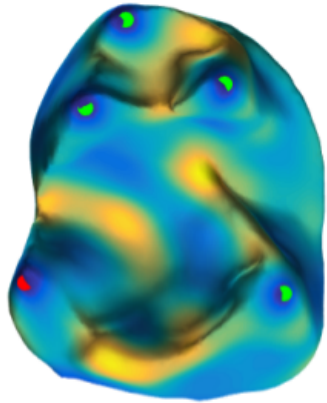
Step 3



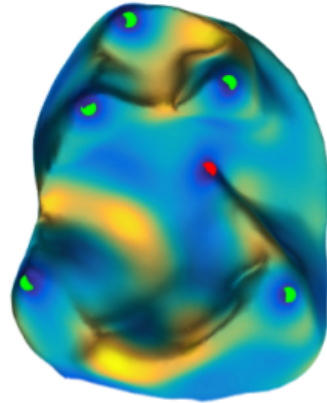
Step 4



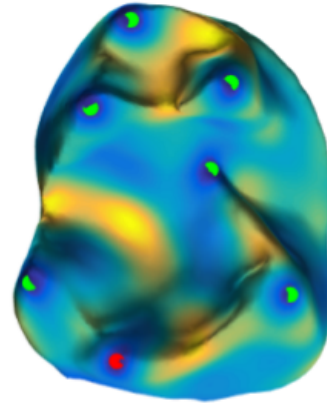
Step 5



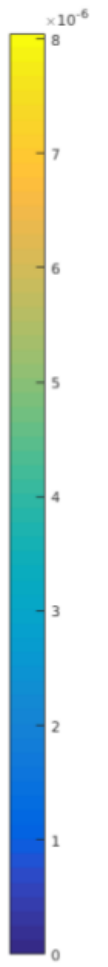
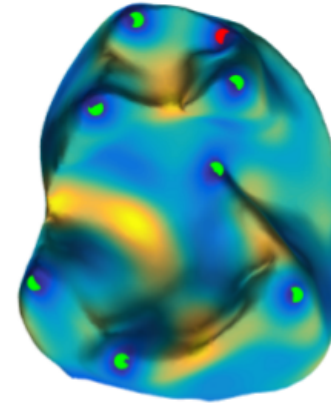
Step 6



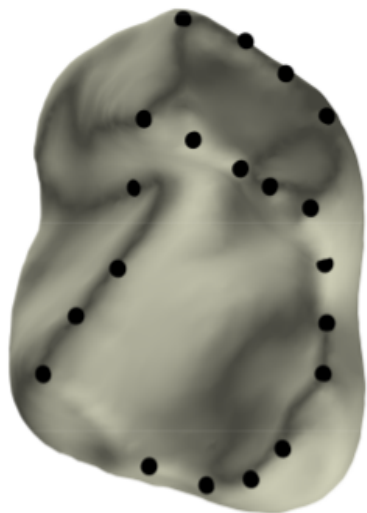
Step 7



Step 8



Gaussian Process Landmarking



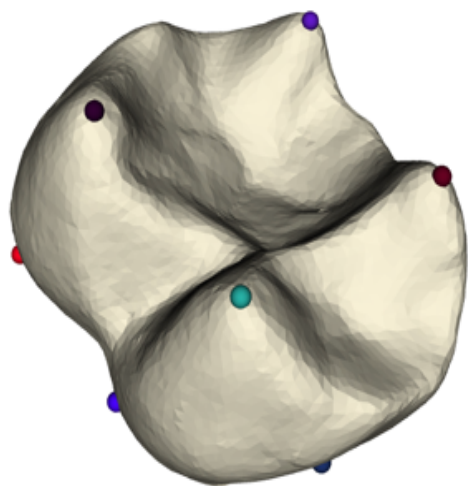
Local Weight Maximum



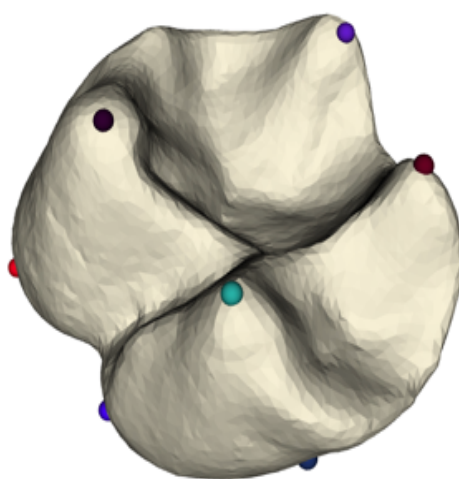
Geodesic Farthest Point Sampling



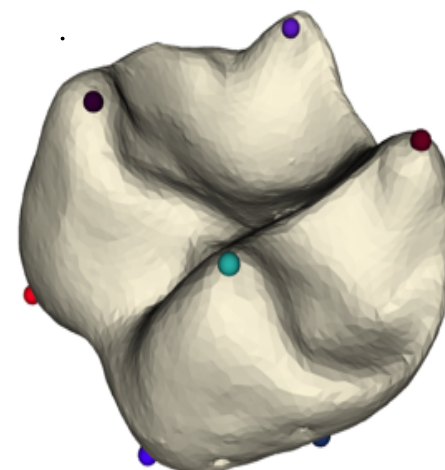
USNM-281751_M759



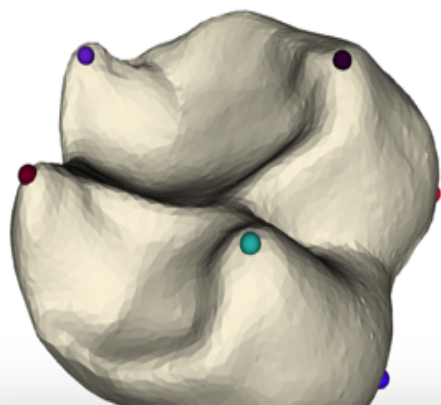
USNM-281758_M761



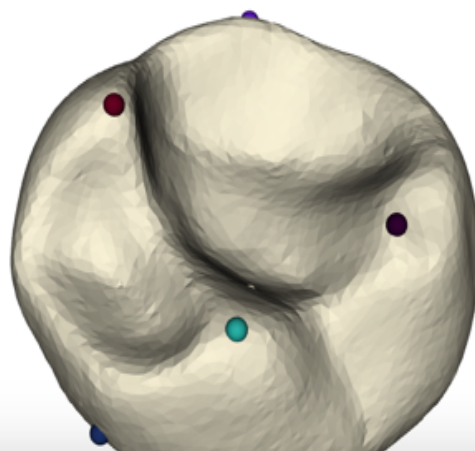
USNM-284782_M1081



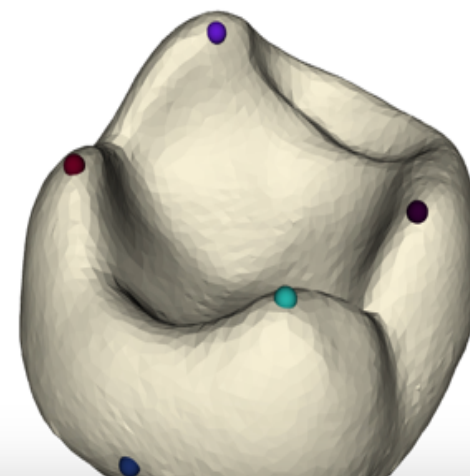
USNM-290601_M1083



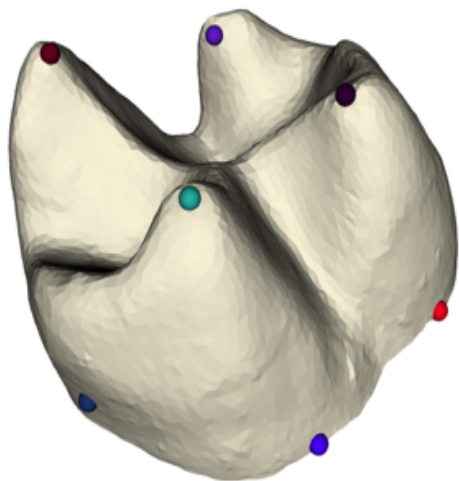
AMNH-
M-67102_M1099



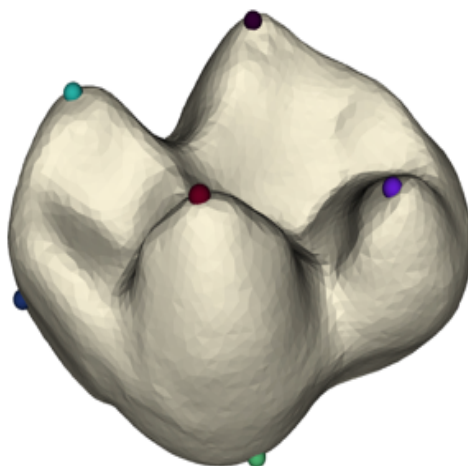
AMNH-
M-71787_M784



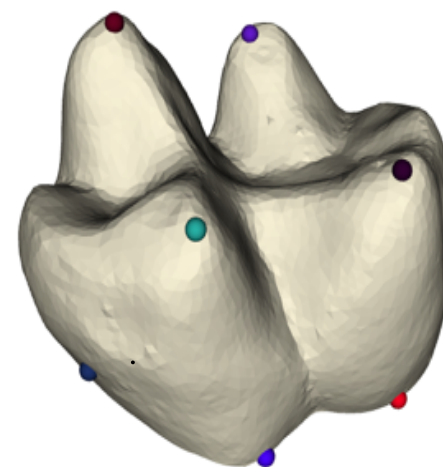
MN-Rio-106_M1214



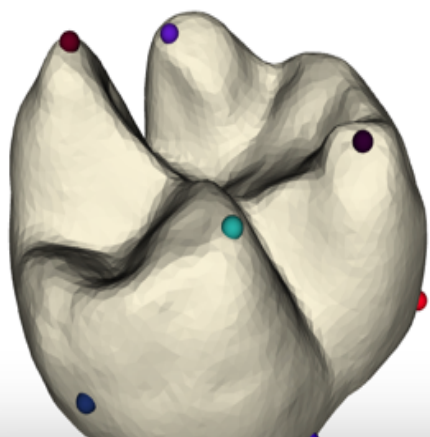
MN-Rio-526_M1205



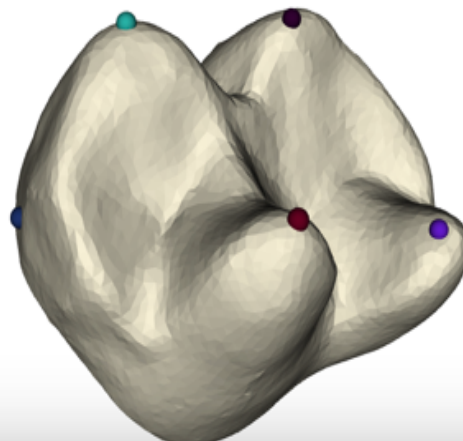
MN-Rio-2718_M1198



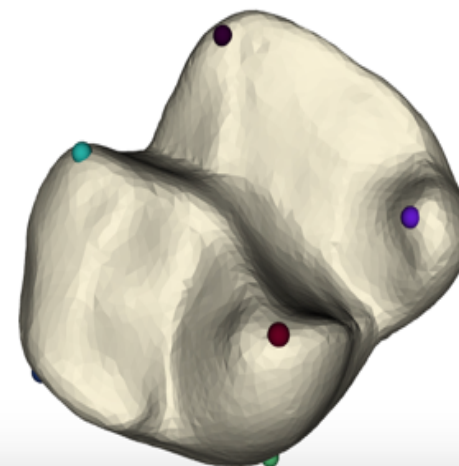
MN-Rio-6699_M1211



MN-
Rio-7724_M1223



MN-
Rio-8513_M1220



Other generalizations in progress:

use a local geometry-adapted version of Procrustes distance.

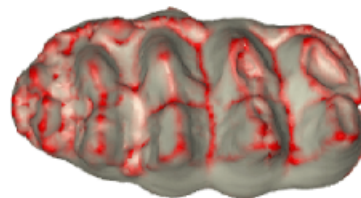
DNE

(AriaDNE)
Shan Shan

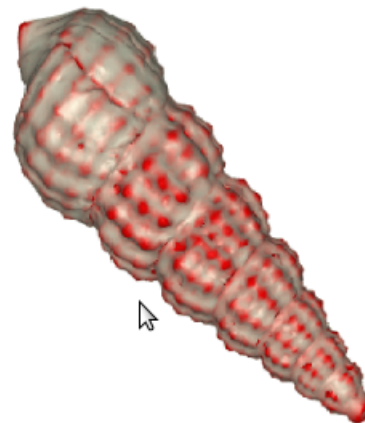
Oryctolagus
0.027



Mammut
0.040



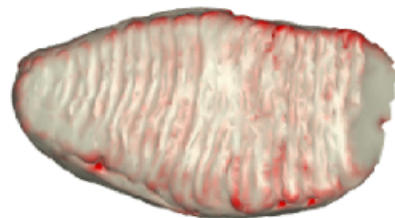
Lirobittium rugatum
0.058



Hemicentetes
0.014



Mammuthus
0.018



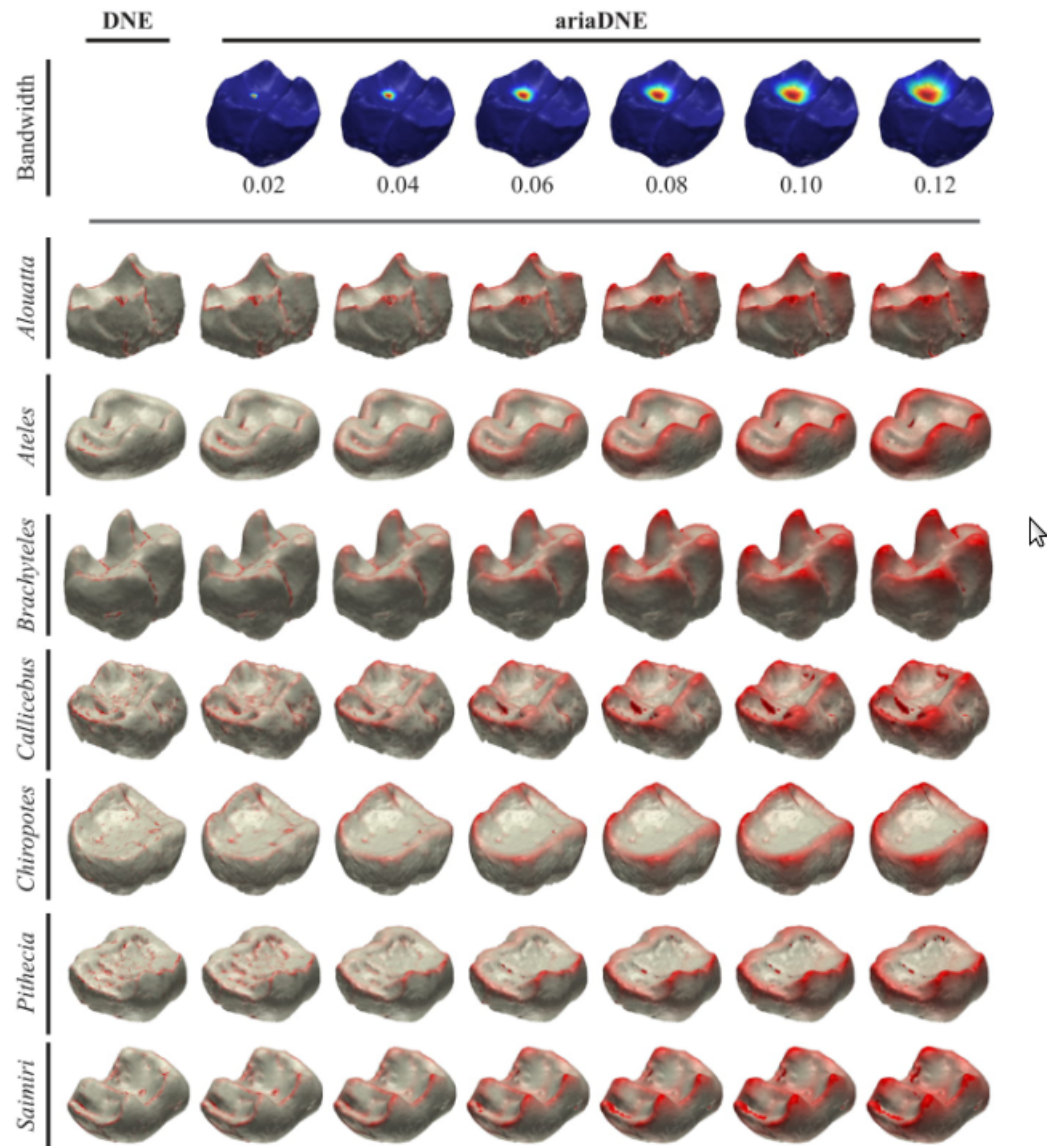
Tornatellaria adelinae
0.009



DNE

(AriaDNE)

Shan Shan



Other generalizations in progress:

- use a local geometry-adapted version of Procrustes distance.

- varying ratios of diffusion times on fiber and base: first optimize registration by emphasizing fiber diffusion; then increasingly emphasize base diffusion to obtain base geometry insights

Happy Birthday, Albert!