

A Growing Self-Organizing Network for Manifold Reconstruction

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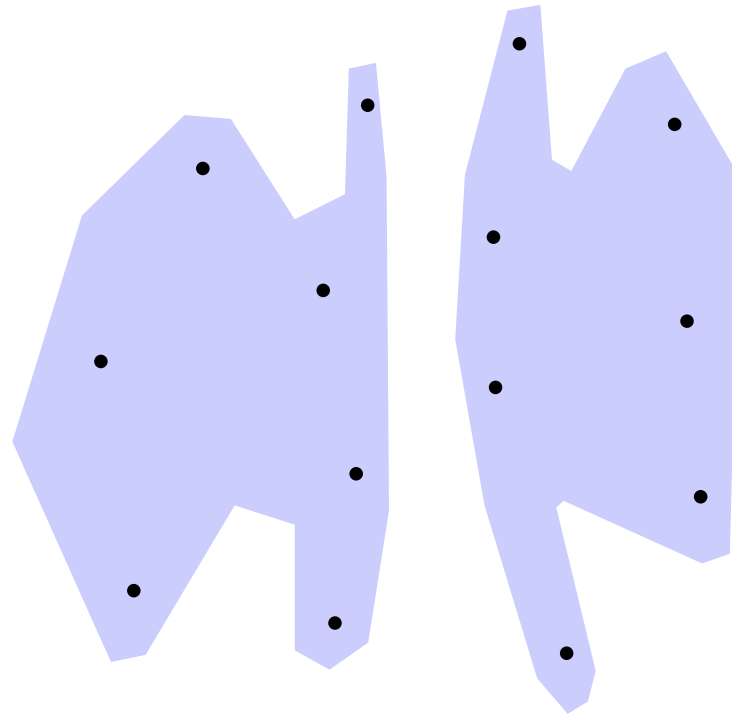
Restricted Delaunay Complex

- **Manifold (a surface embedded in \mathbf{R}^2)**



Restricted Delaunay Complex

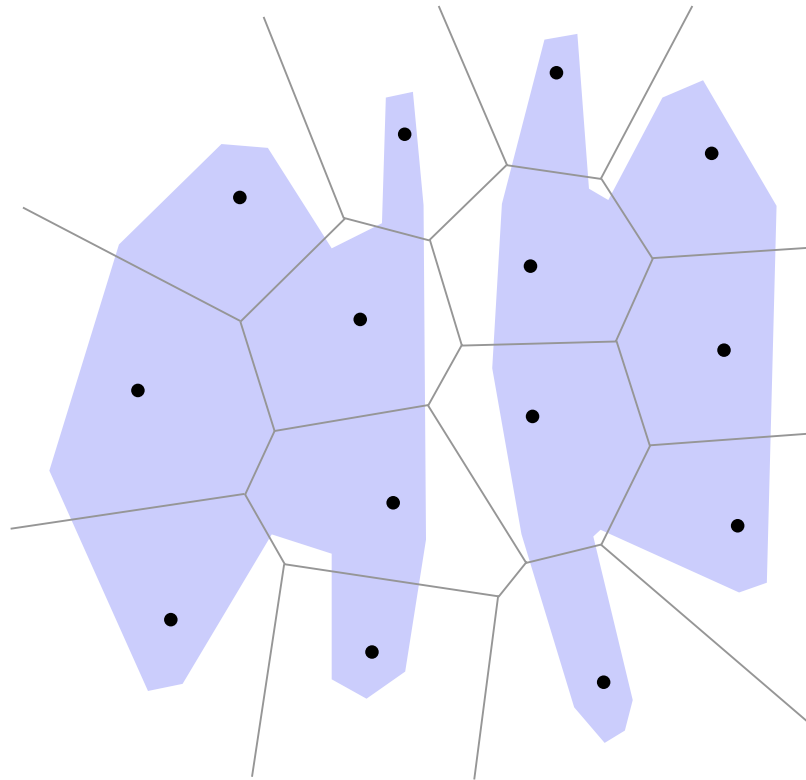
- Point sample (*landmarks*) of the manifold



Restricted Delaunay Complex

- **Voronoi complex of the landmarks**

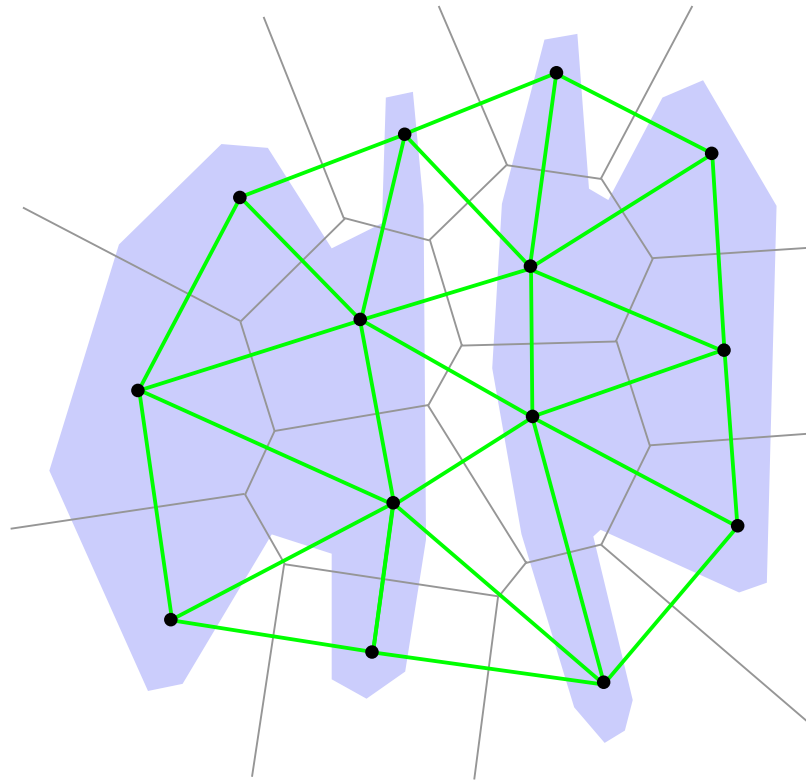
Each cell contains all points of \mathbf{R}^2 being closer to a specific *landmark*



Restricted Delaunay Complex

- **Delaunay graph of the landmarks**

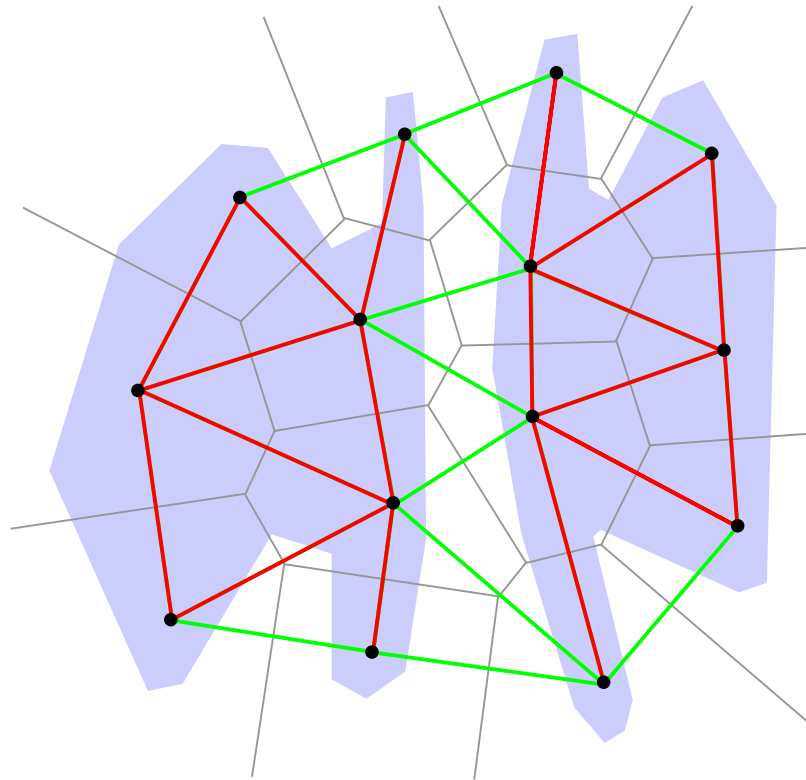
An edge connects each two landmarks whose Voronoi cells have a common *boundary*



Restricted Delaunay Complex

- **Restricted Delaunay graph of the landmarks**

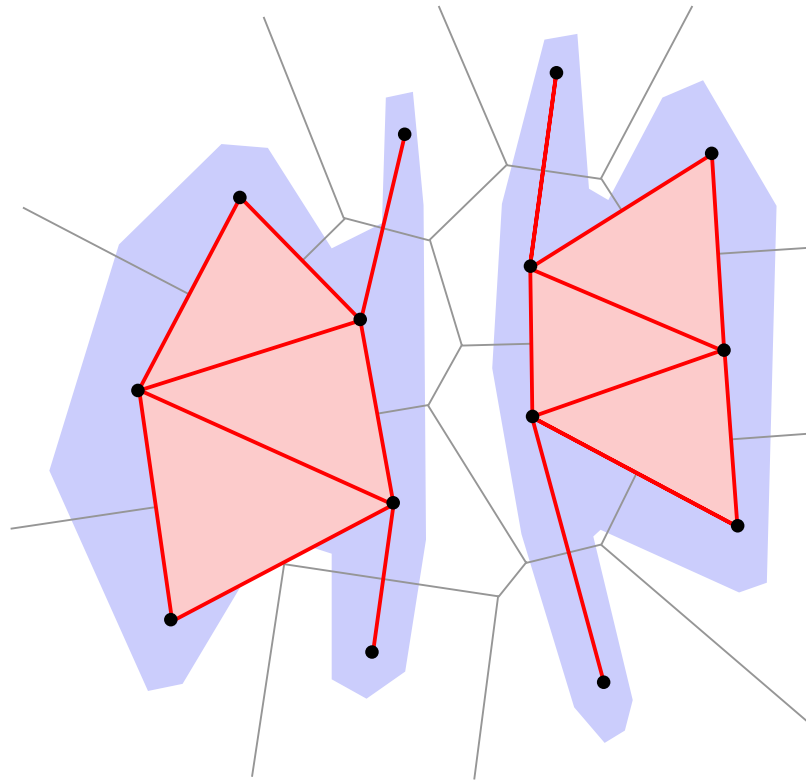
An edge connects each two landmarks whose Voronoi cells have a common *boundary* which intersects the manifold M



Restricted Delaunay Complex

- **Restricted Delaunay complex of the landmarks**

A $(n - 1)$ -dimensional n -face corresponds to n landmarks whose Voronoi cells have a common *boundary* which intersects M

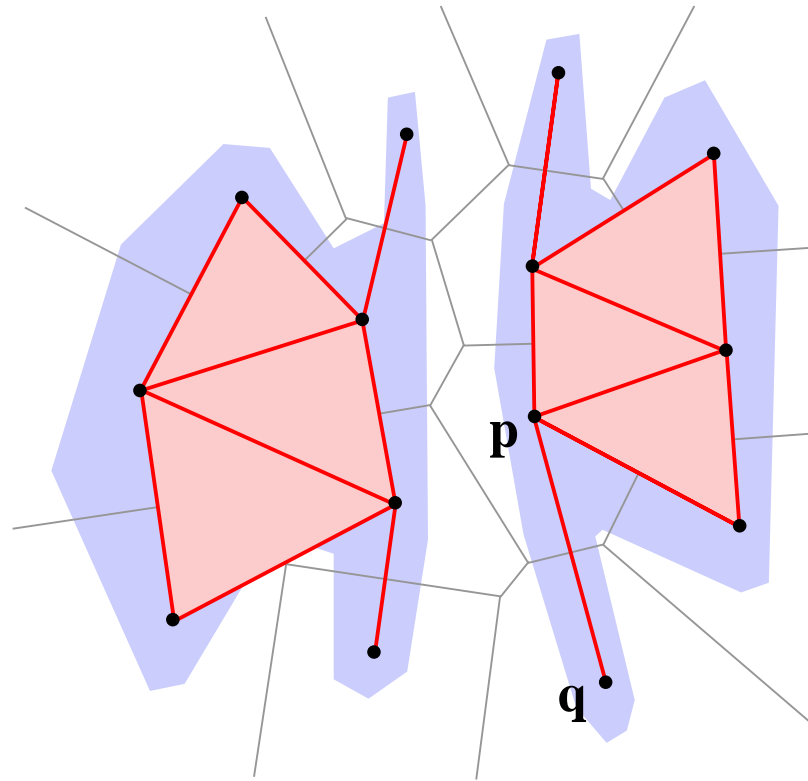


Restricted Delaunay Complex

- **Restricted Delaunay complex of the landmarks**

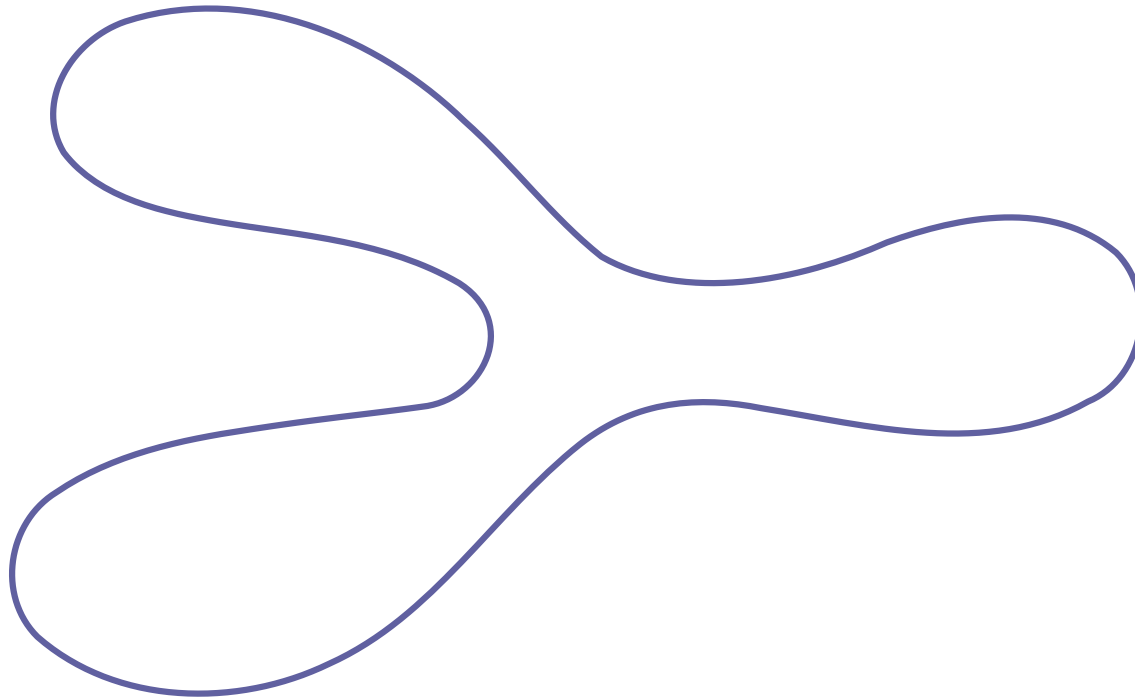
The complex, in general, is *not* homeomorphic to the manifold

Here, for instance, the neighborhoods of either **p** or **q** have no counterparts in M



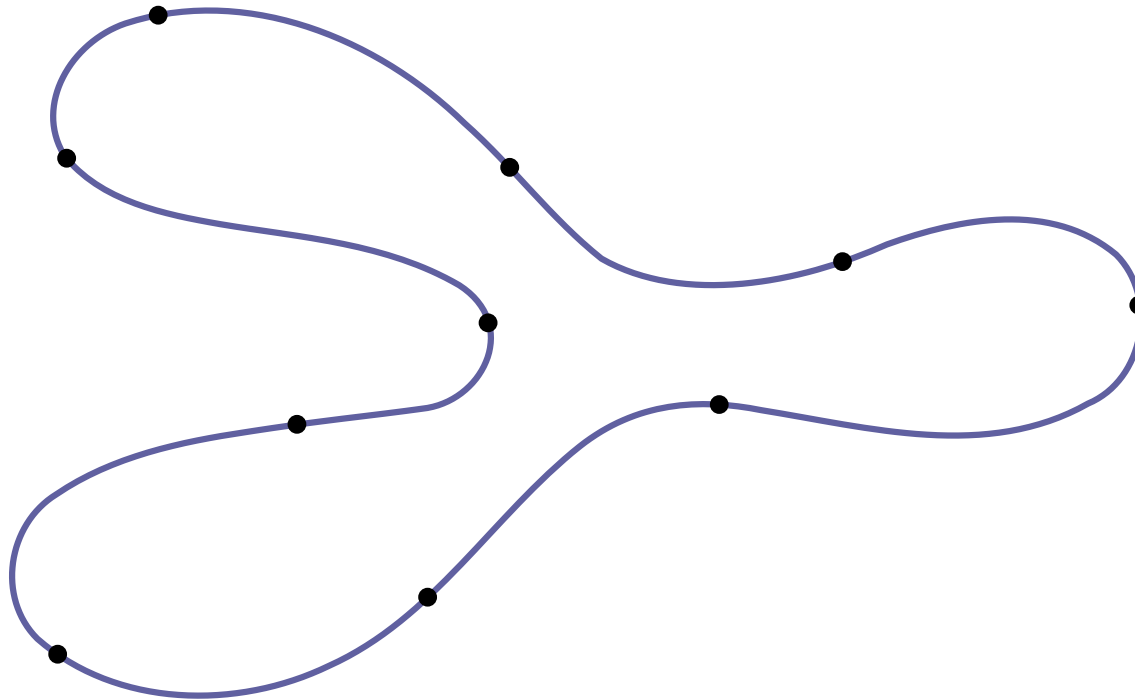
Restricted Delaunay Complex and Homeomorphism

- **Manifold (a curve embedded in \mathbf{R}^2)**



Restricted Delaunay Complex and Homeomorphism

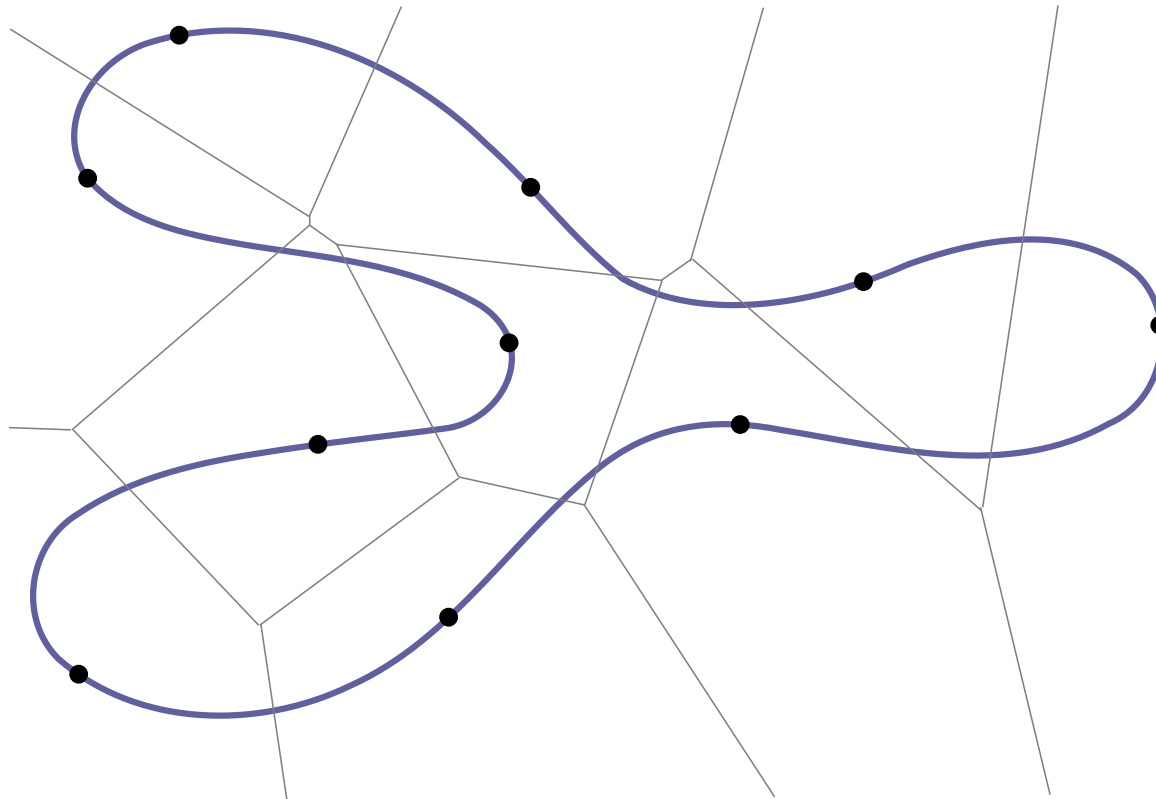
- A first point sample (*landmarks*) of the manifold



Restricted Delaunay Complex and Homeomorphism

- **Voronoi complex**

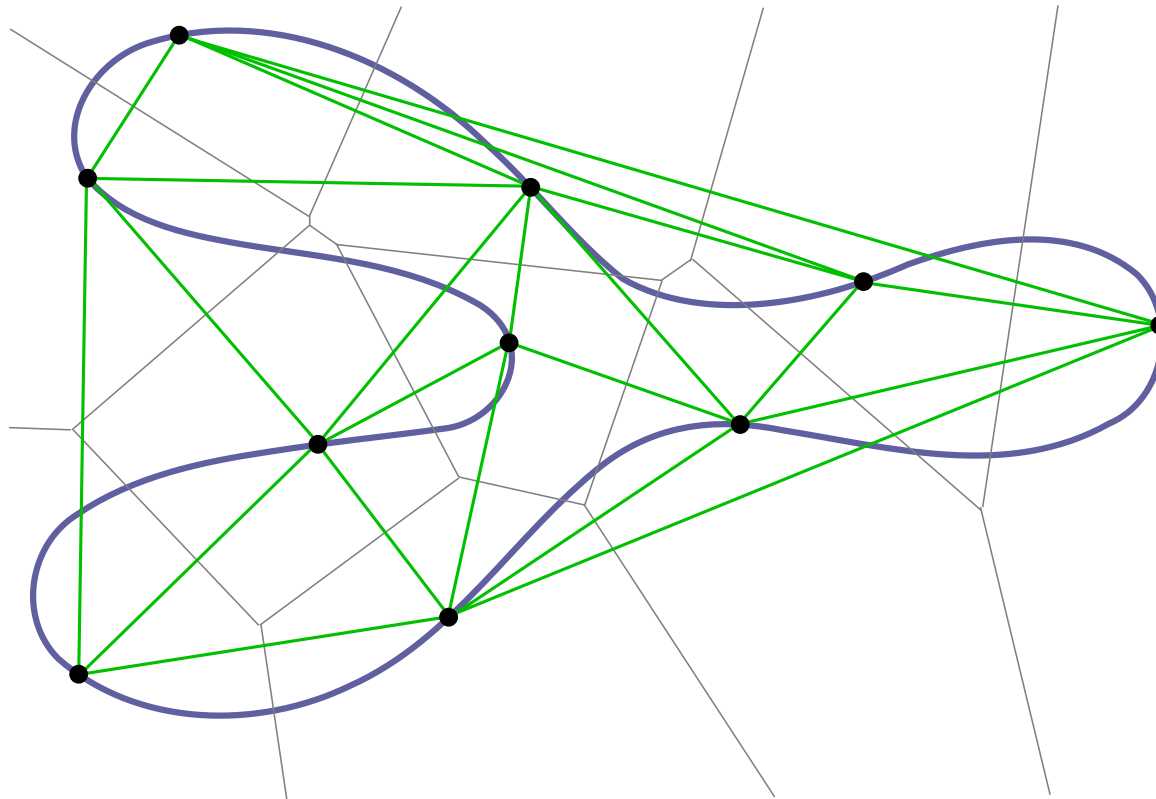
Each cell contains all points of \mathbf{R}^2 being closer to a specific *landmark*



Restricted Delaunay Complex and Homeomorphism

■ Delaunay graph

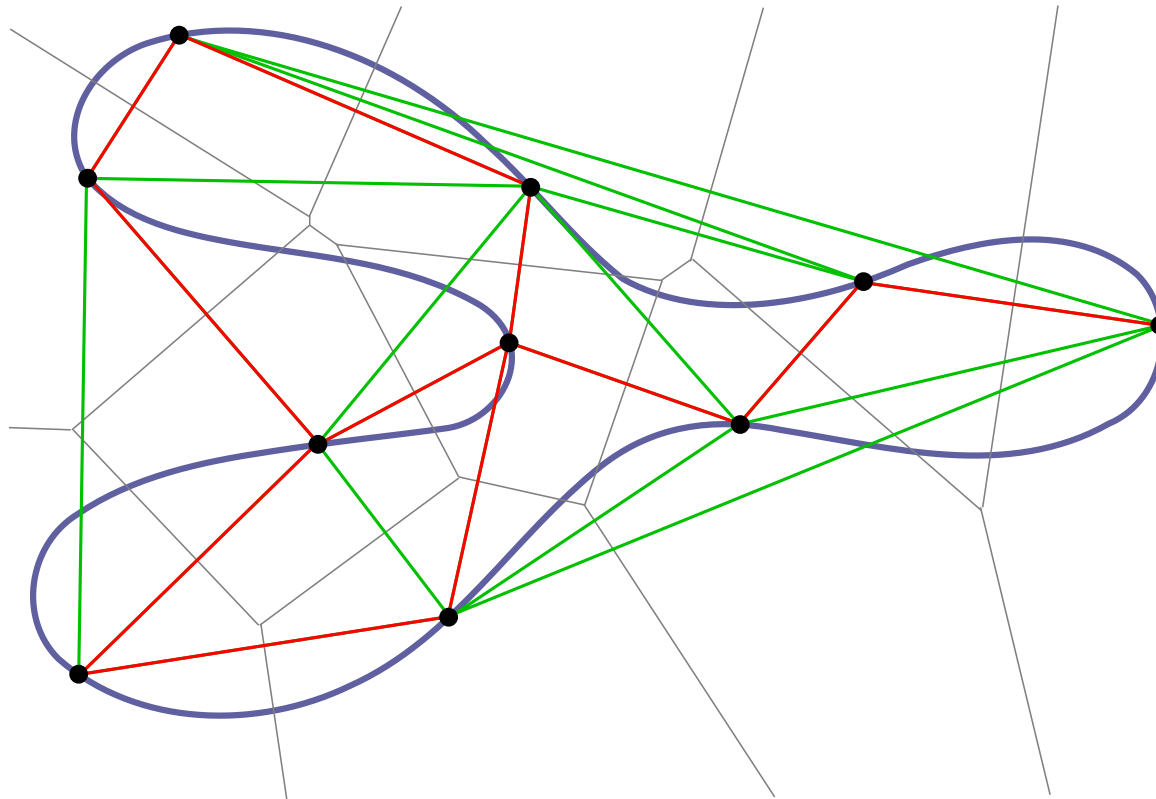
An edge connects each two landmarks whose Voronoi cells have a common *boundary*



Restricted Delaunay Complex and Homeomorphism

■ Restricted Delaunay graph

An edge connects each two landmarks whose Voronoi cells have a common *boundary* which intersects M

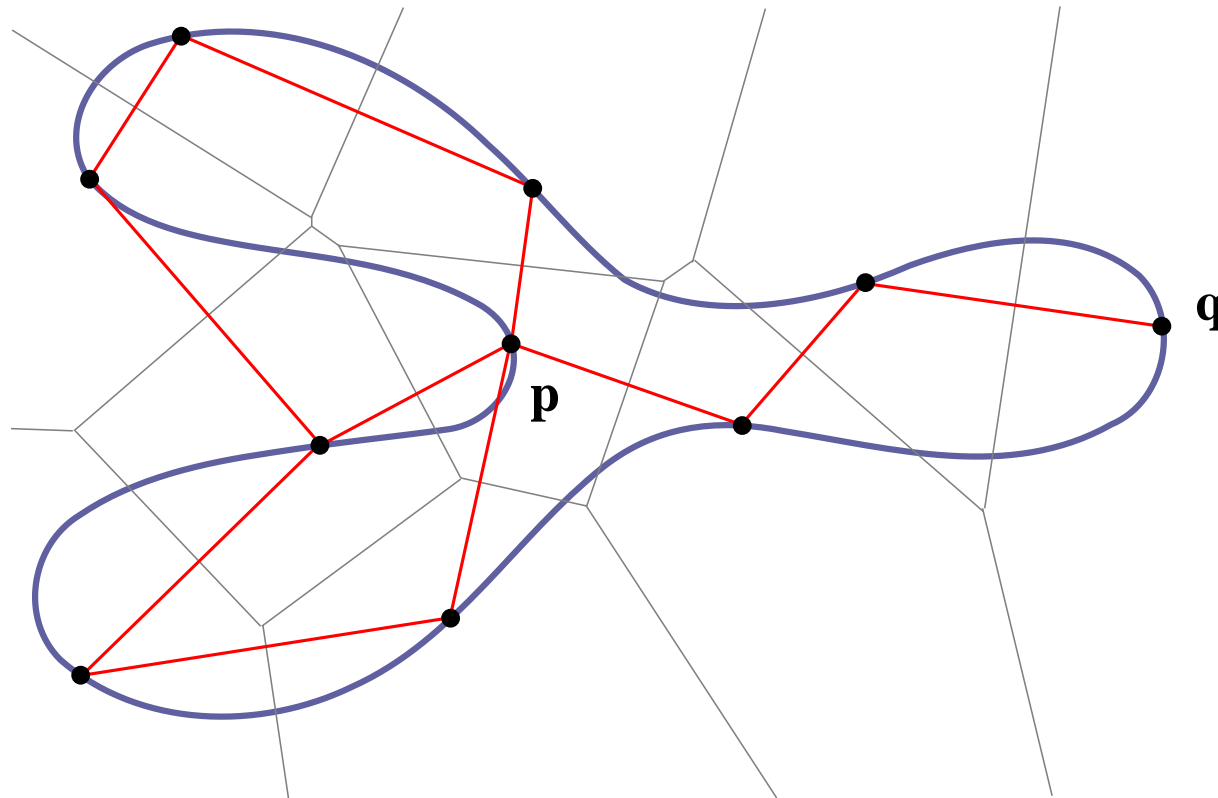


Restricted Delaunay Complex and Homeomorphism

■ Restricted Delaunay graph

Once again and in general, the complex is *not* homeomorphic to the manifold

Here, for instance, the neighborhoods of either p or q have no counterparts in M

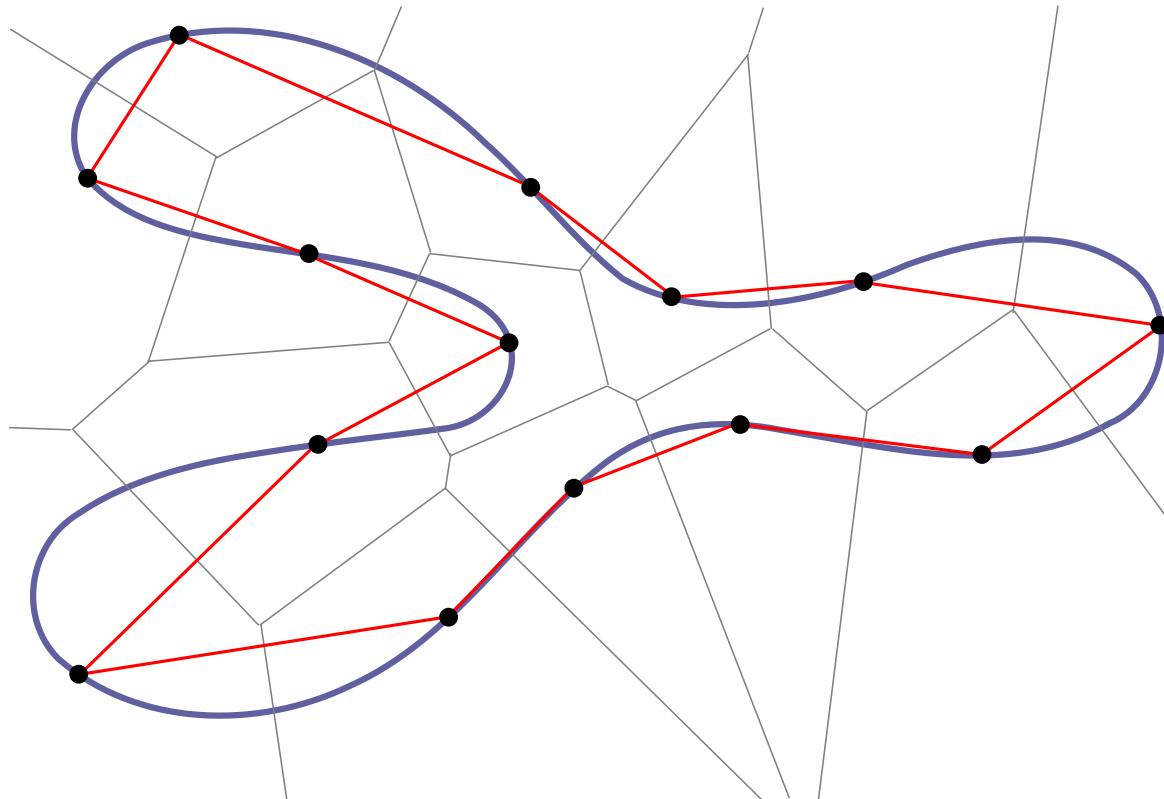


Restricted Delaunay Complex and Homeomorphism

- **Want homeomorphism?**

Just add more landmarks.

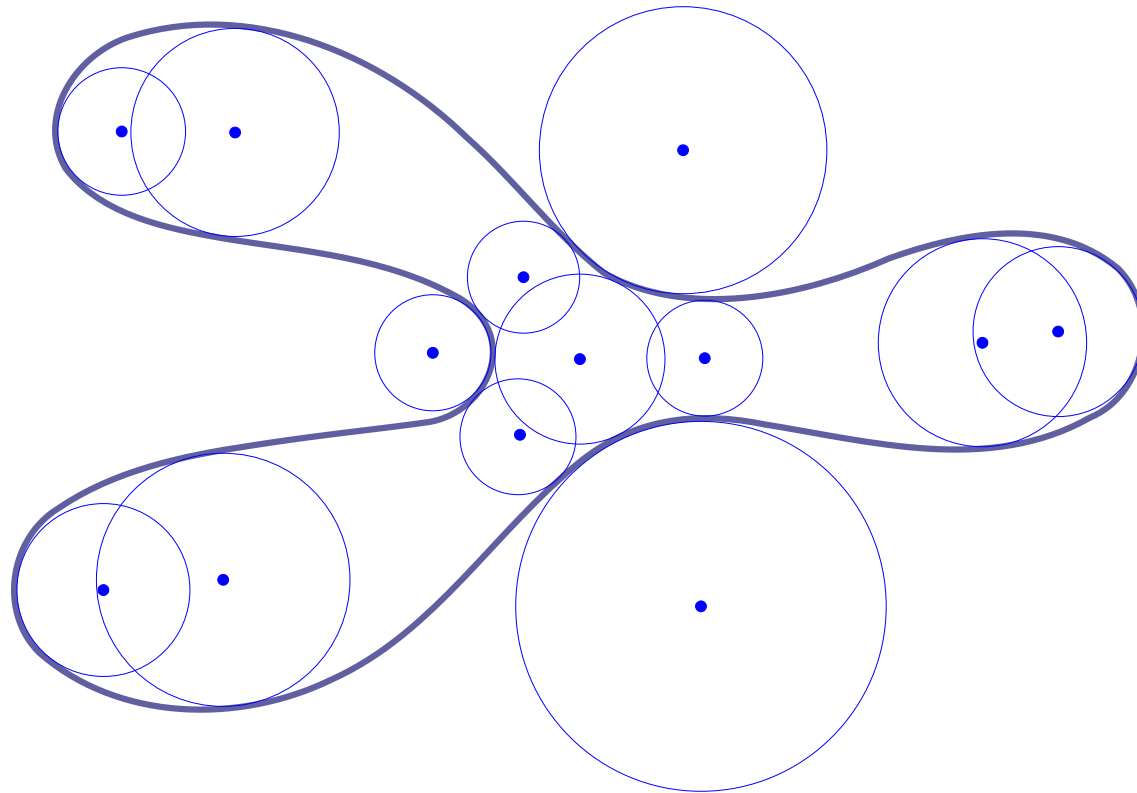
(There exists a *density threshold*)



ε -sample

- **Medial balls**

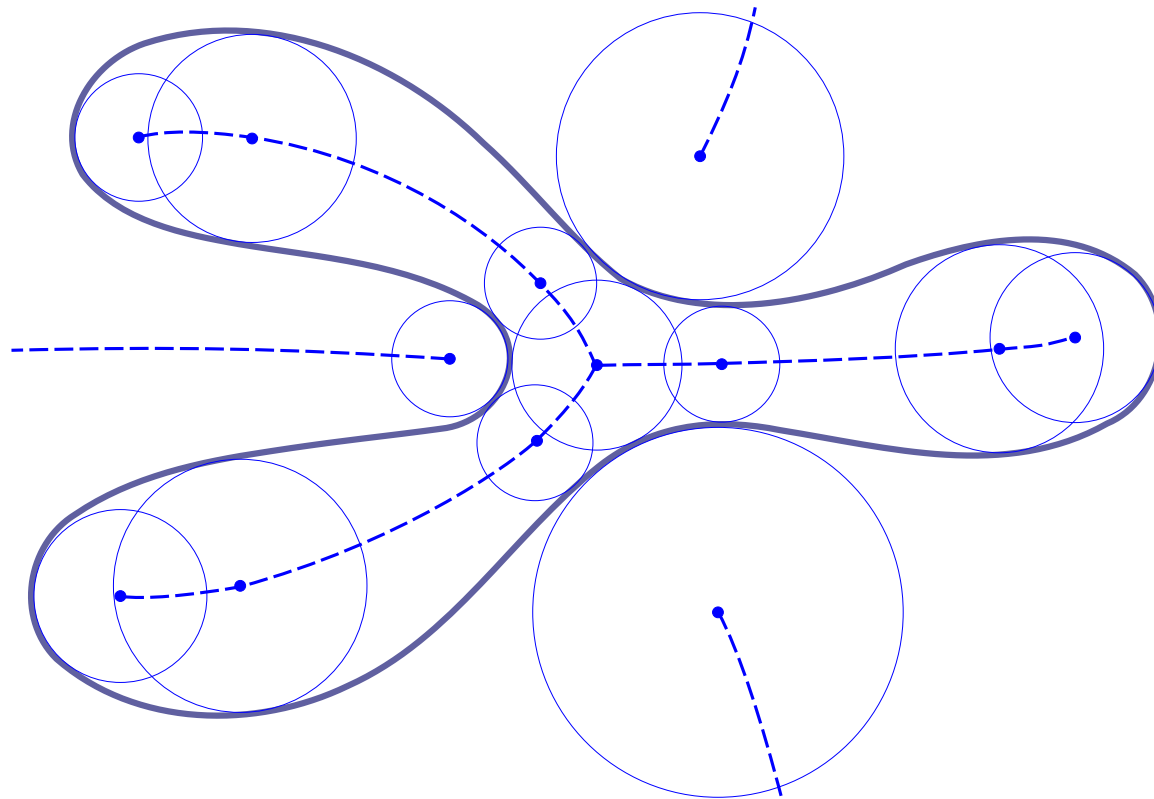
Maximal balls whose interiors are empty of any points from M



ε -sample

- **Medial axis**

The closure of the set of points that are centers of maximal balls

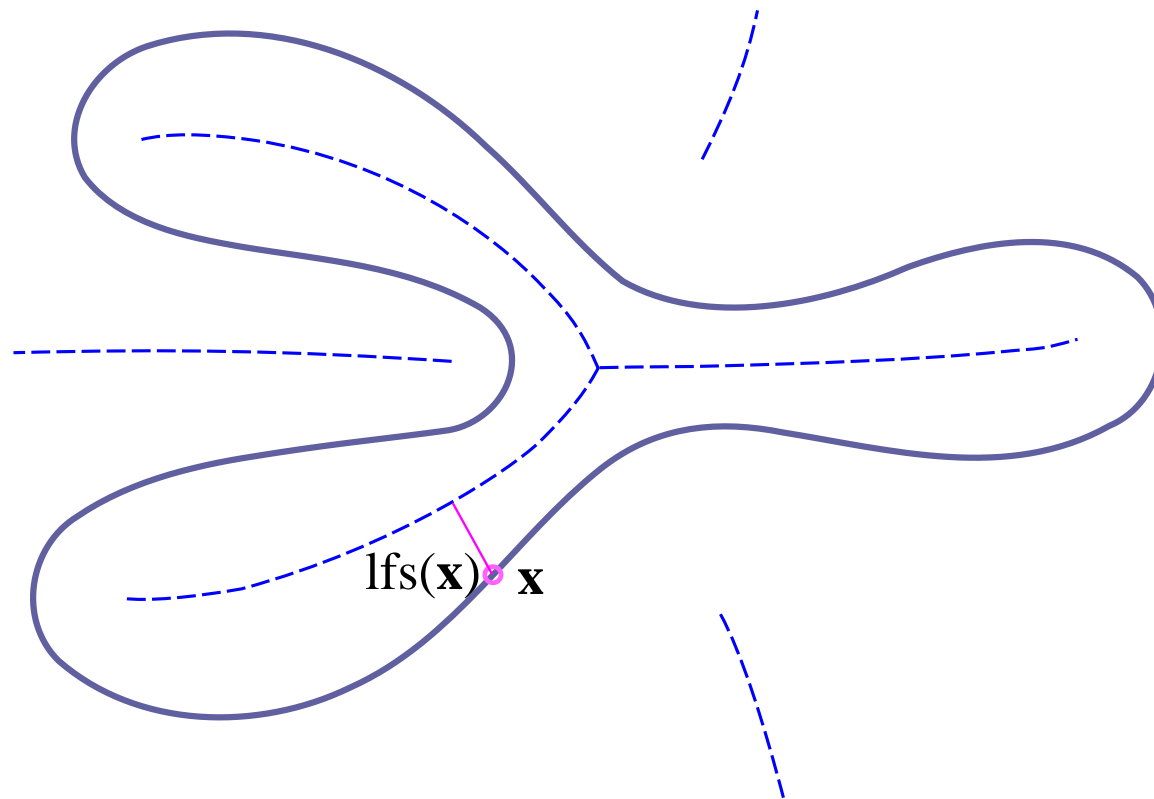


ε -sample

▪ Local Feature Size

(at a point \mathbf{x} on M)

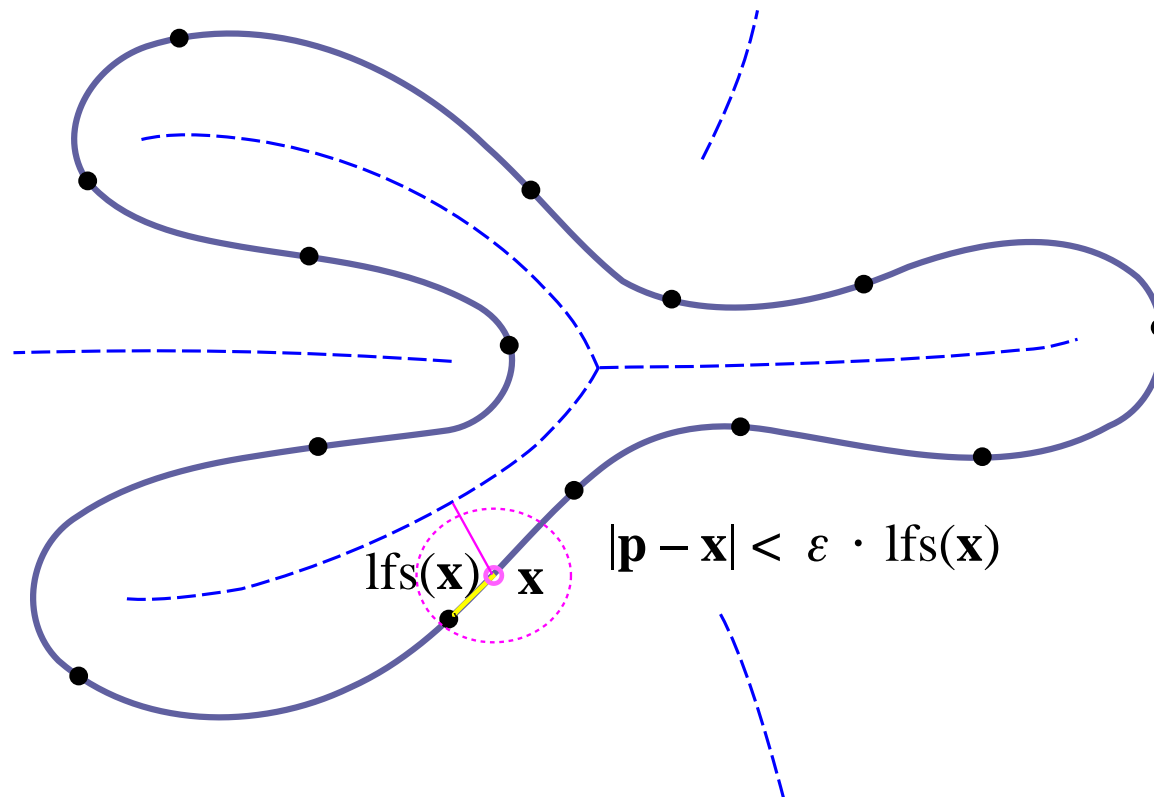
It is the distance between \mathbf{x} and the medial axis



ε -sample

■ ε -sample

A set of landmarks such that every point \mathbf{x} on M is at most $\varepsilon \cdot \text{lfs}(\mathbf{x})$ away from the closest landmark \mathbf{p}

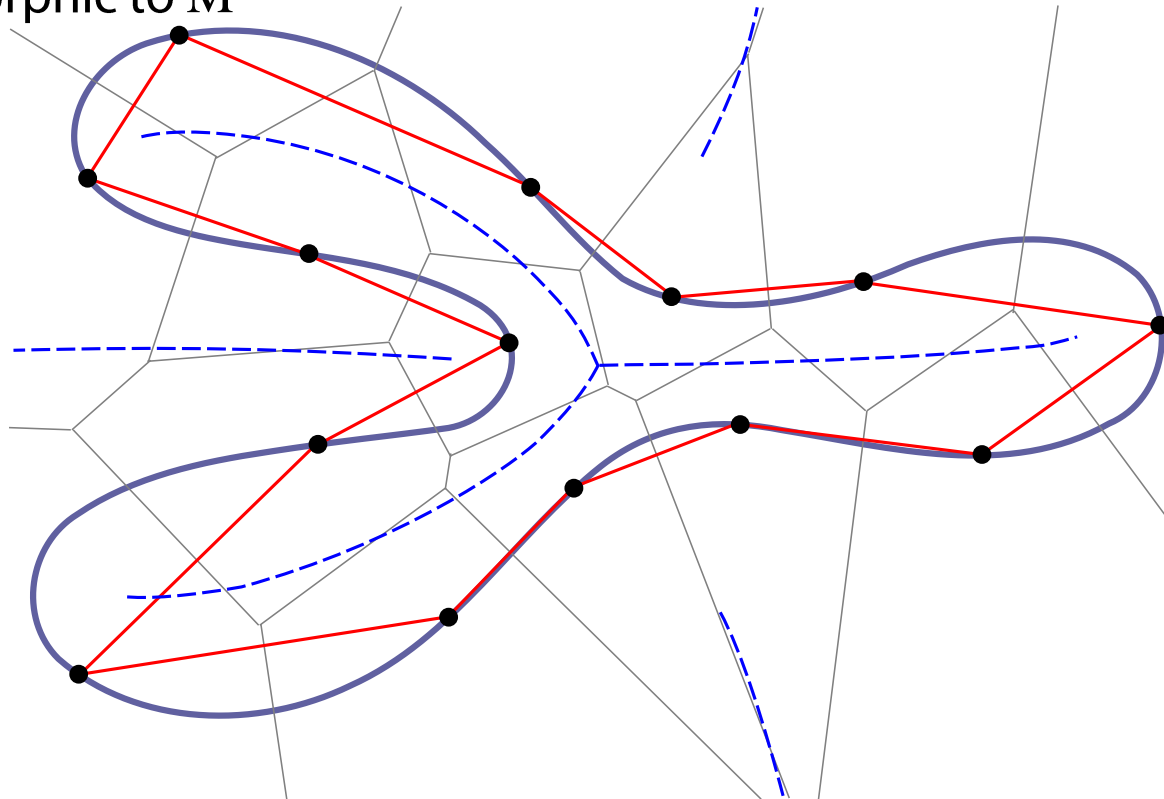


ε -sample

■ ε -sample and homeomorphism

[Amenta et al., 2000]

If M is compact, closed and *smooth*, there exists a positive ε such that the restricted Delaunay complex for any ε -sample of M is homeomorphic to M



ε -sample

▪ **The restricted Delaunay complex of an ε -sample**

When M is compact, closed and *smooth* and ε is sufficiently small

- It is homeomorphic to M
- The Hausdorff distance to M is $O(\varepsilon^2)$
- It allows a reliable estimate of curvatures, normals, lengths or areas of M

▪ **Limitations**

It works only with manifolds of dimension 1 or 2

Although the dimension of the ambient space could be any

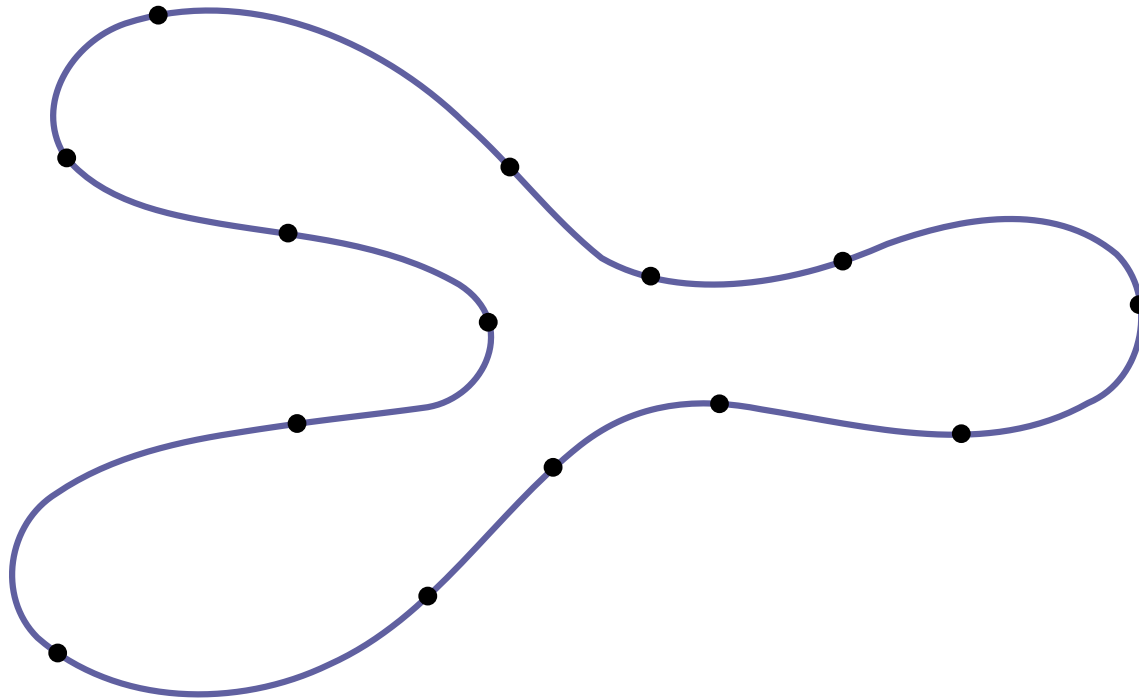
[Oudot, 2008]

For manifolds of dimension greater than 2,
no positive value of ε guarantees
that an ε -sample has the properties above

A weighted Delaunay complex could bring those properties back (but this is another story)

Witness complex

- **How can the restricted Delaunay complex be constructed?**
(From a given set of landmarks)

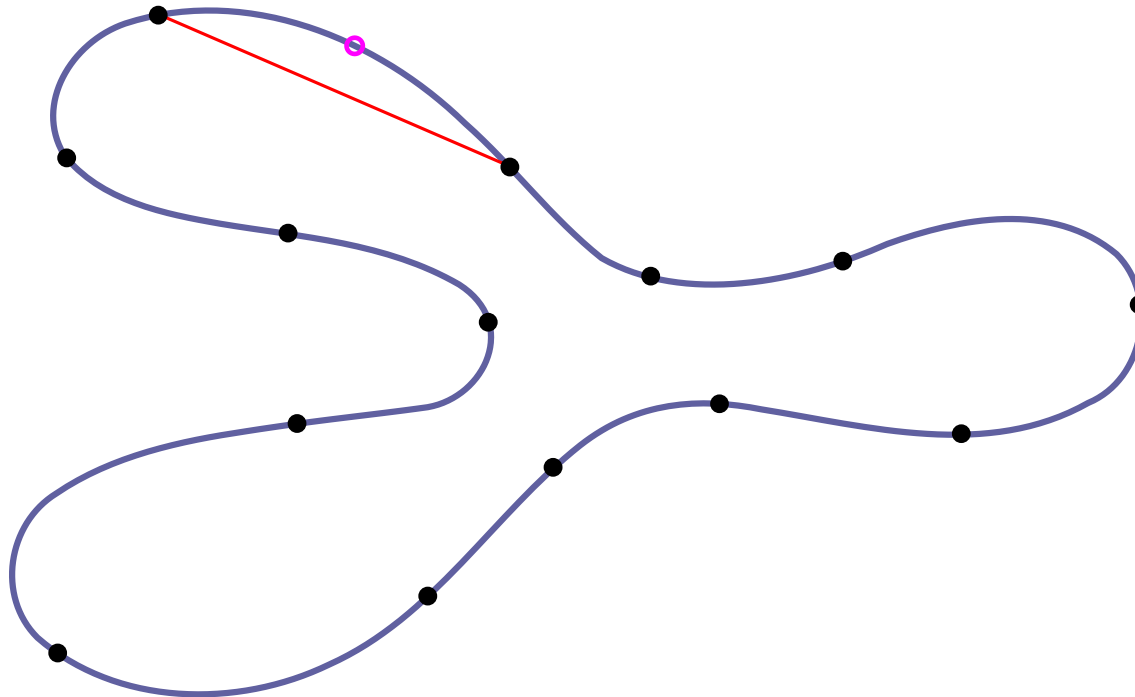


Witness complex

- **Try sampling the manifold at random**

For each sample, add a connection between the two closest landmarks

The sampled point is deemed a *witness* for the corresponding connection

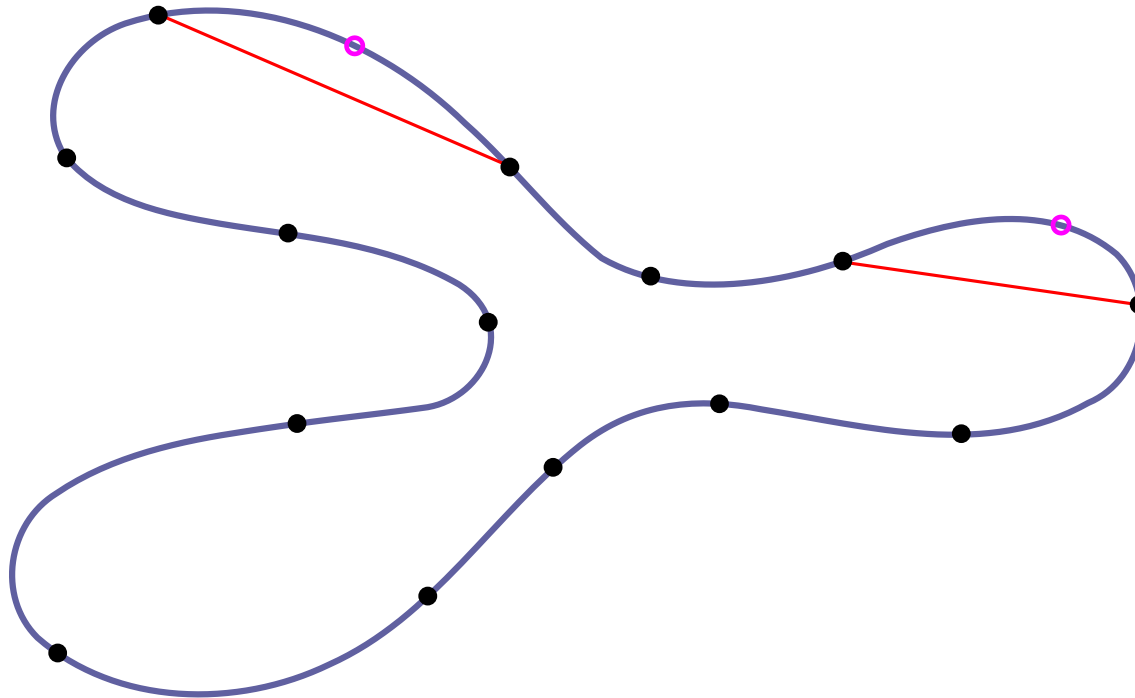


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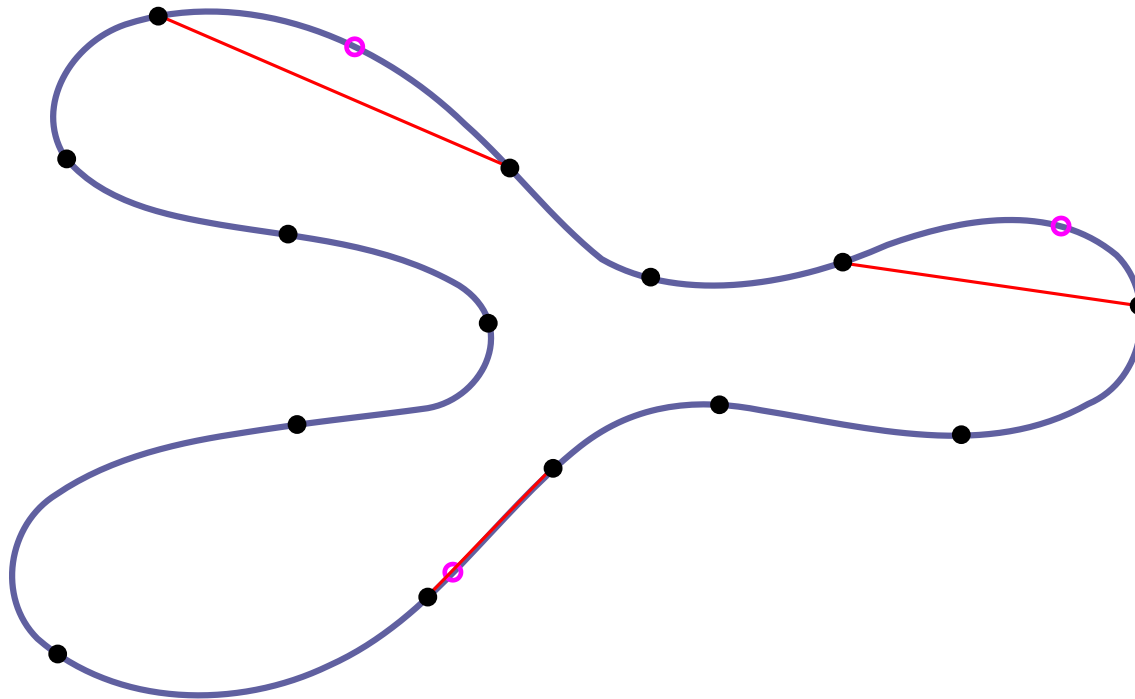


Witness complex

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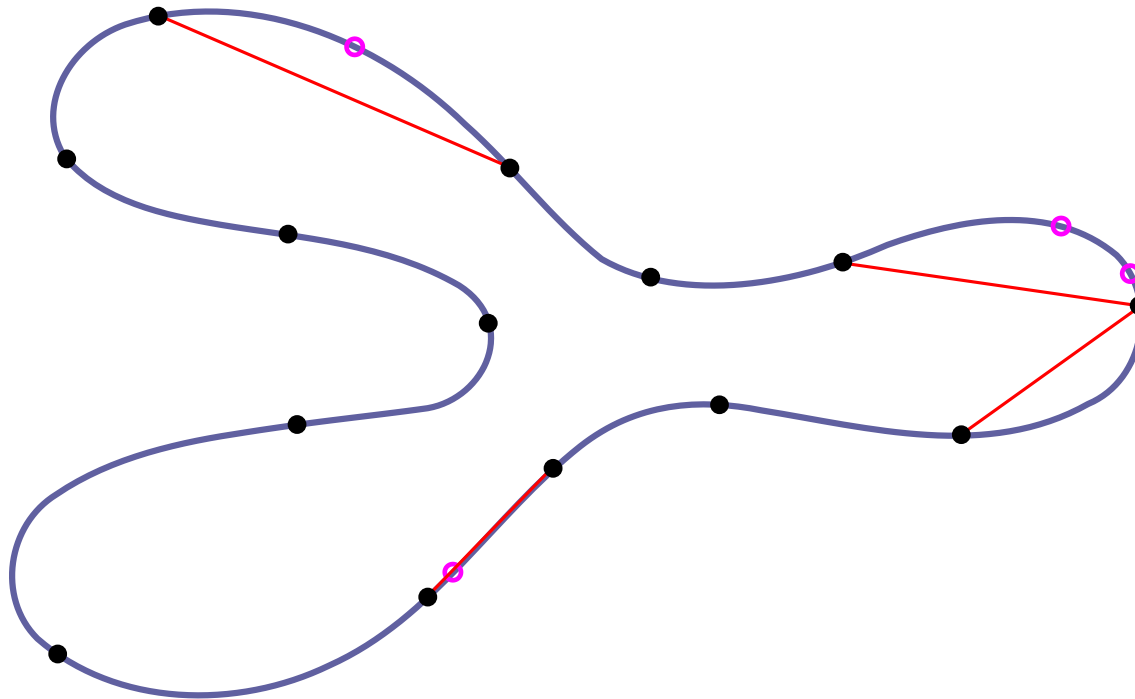


Witness complex

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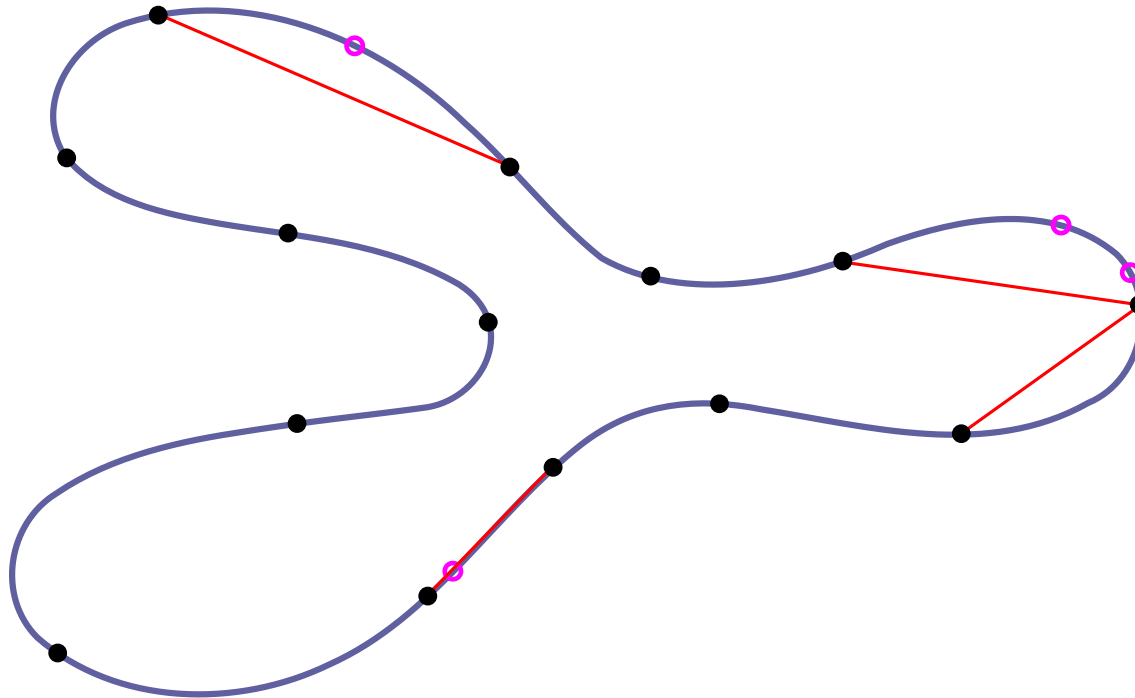
The sampled point is deemed a *witness* for the corresponding connection



Witness complex

■ Witness complex

It is the structure obtained by taking the sampling process to the limit
i.e. when the whole M has been sampled

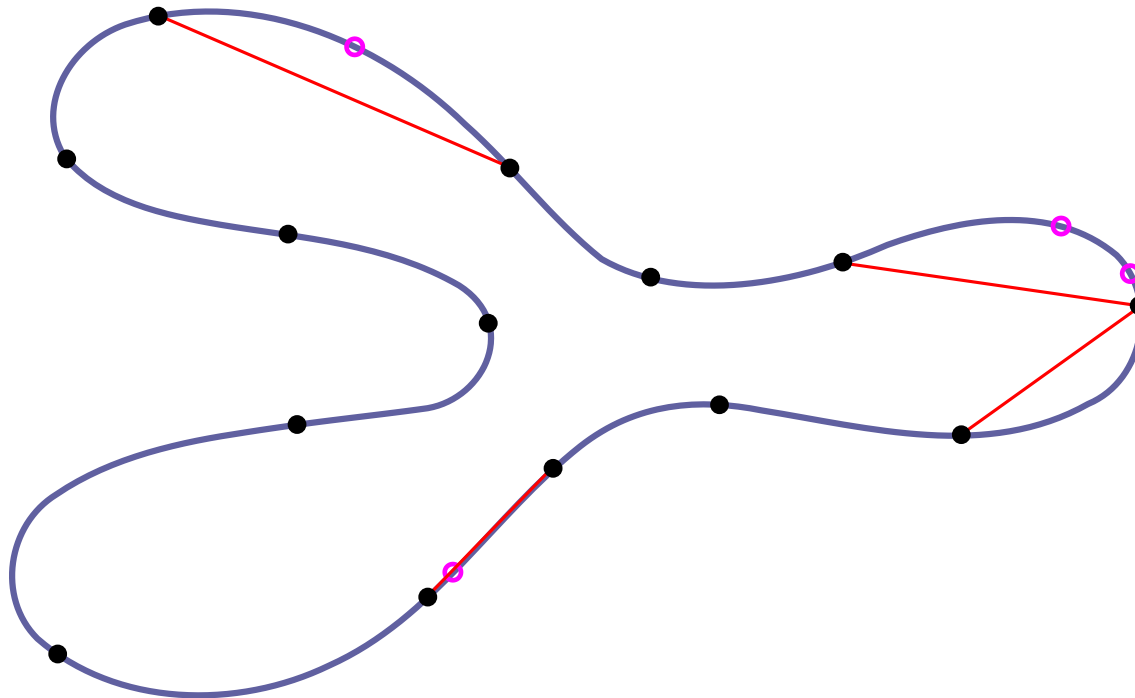


Witness complex

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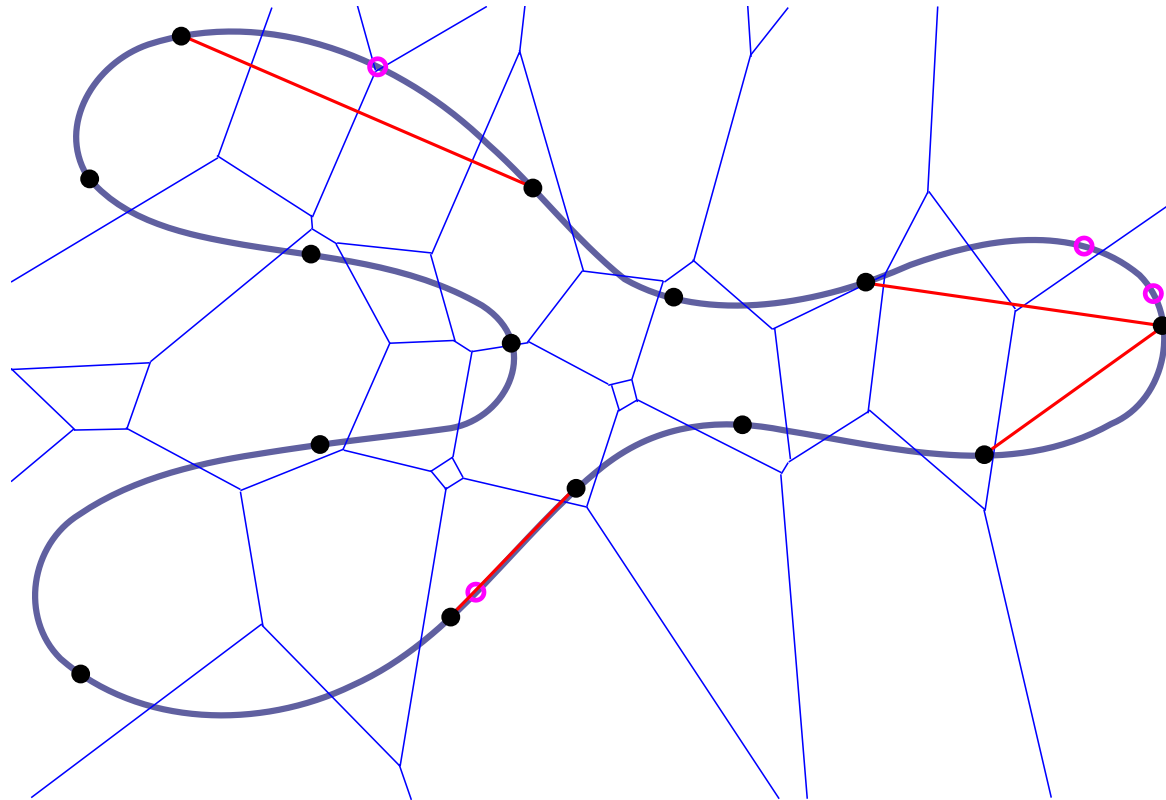
Will it coincide with the restricted Delaunay complex?



Witness complex

- **Second-order Voronoi complex**

Each cell contains all points of \mathbf{R}^2 being closer to a specific pair of landmarks

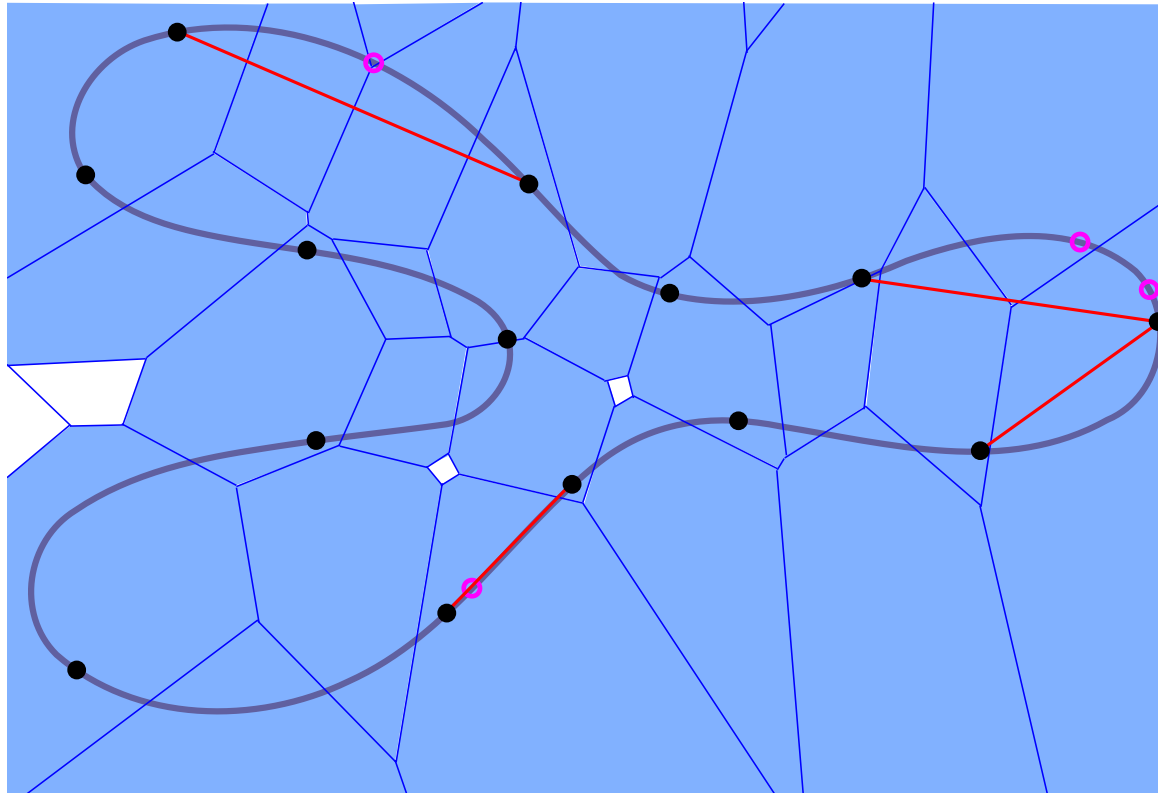


Witness complex

- **Second-order Voronoi complex**

Each cell contains all points of \mathbf{R}^2 being closer to a specific pair of landmarks

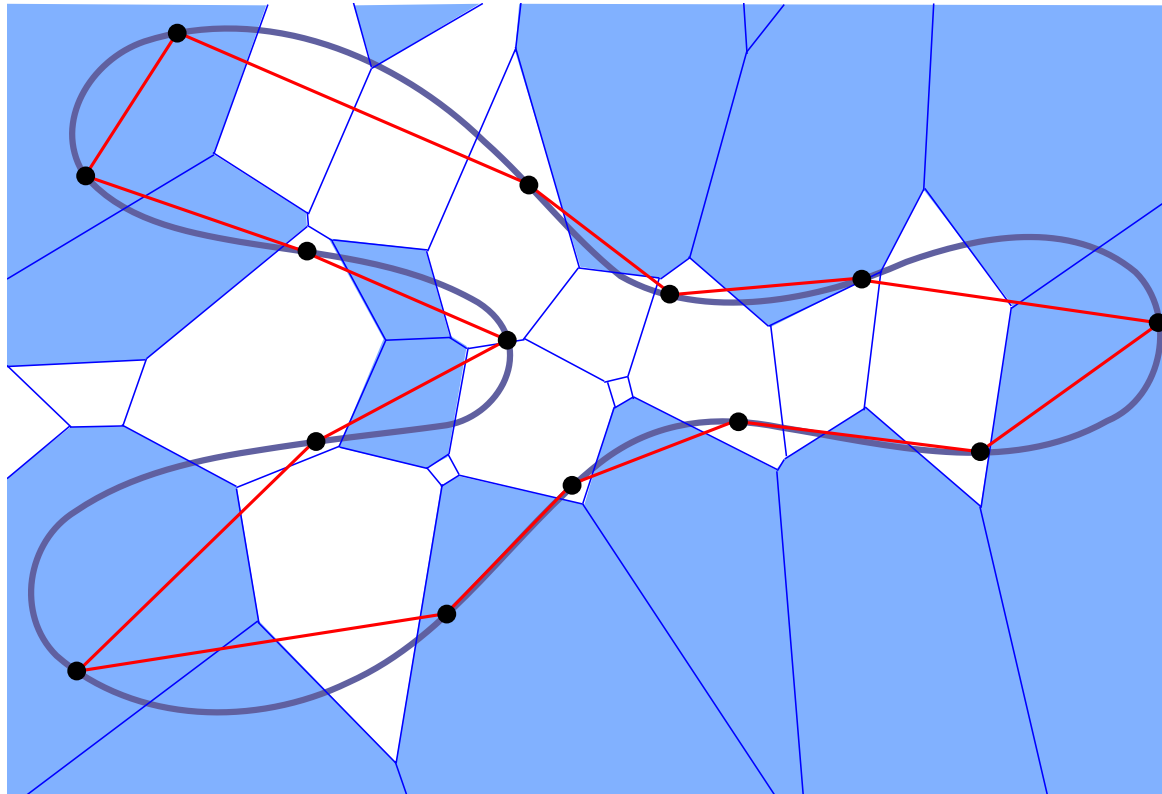
Therefore, each cell intersecting M contains *witnesses* for one connection



Witness complex

- **Second-order Voronoi complex and witness complex**

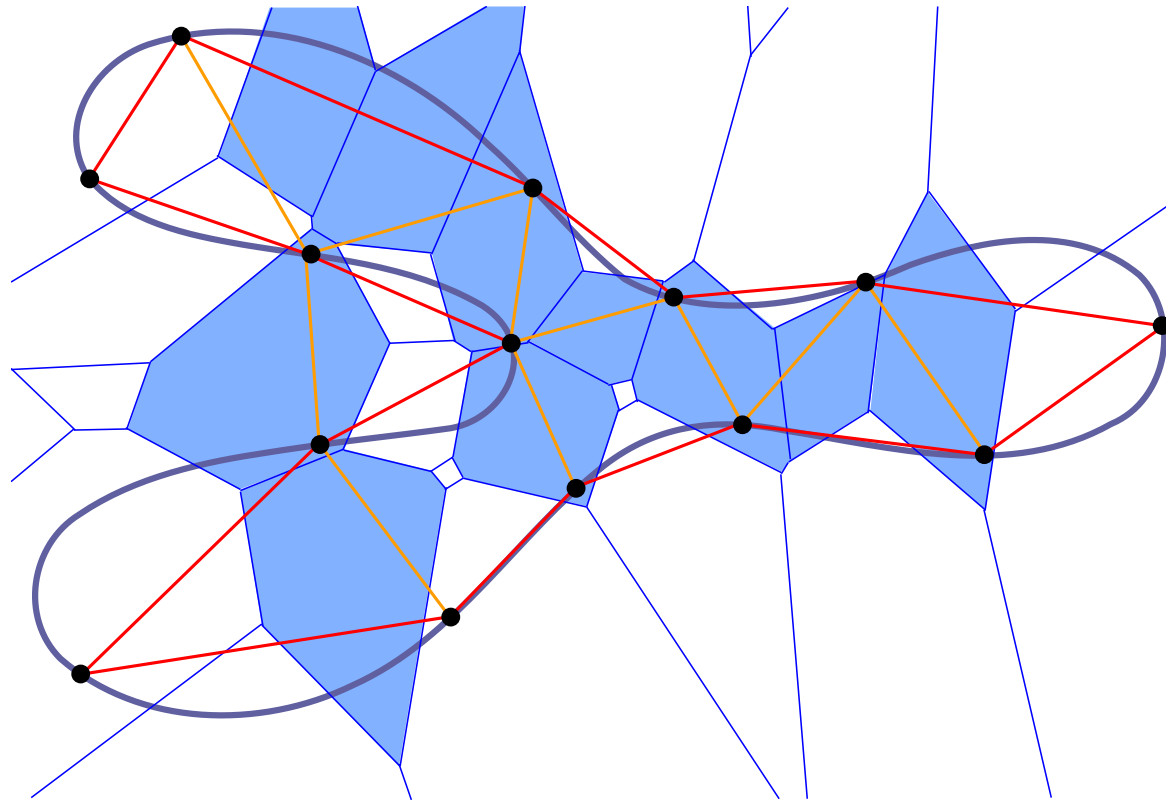
Certainly, there are witnesses for the restricted Delaunay complex



Witness complex

- **Second-order Voronoi complex and witness complex**

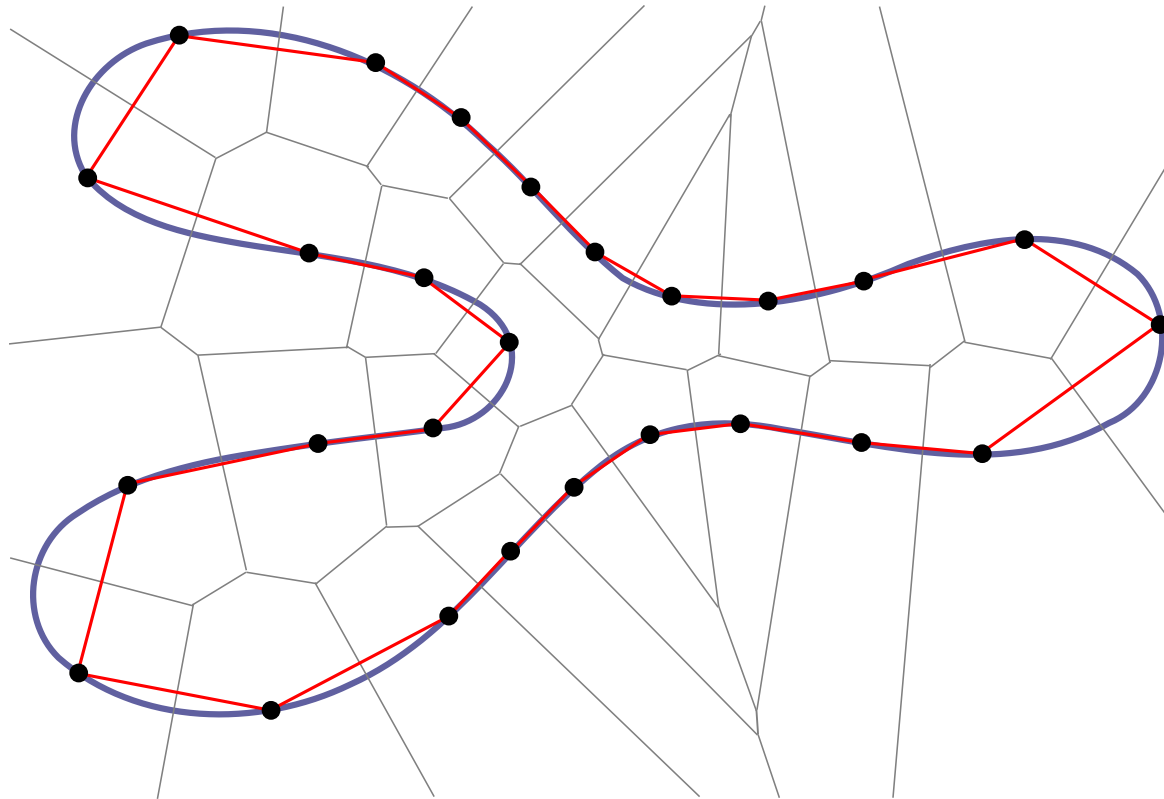
Certainly, there are witnesses for the restricted Delaunay complex but there will be also witnesses for a few extra connections ...



Witness complex

- **Witness complex and the restricted Delaunay complex**

The solution? Add even more landmarks

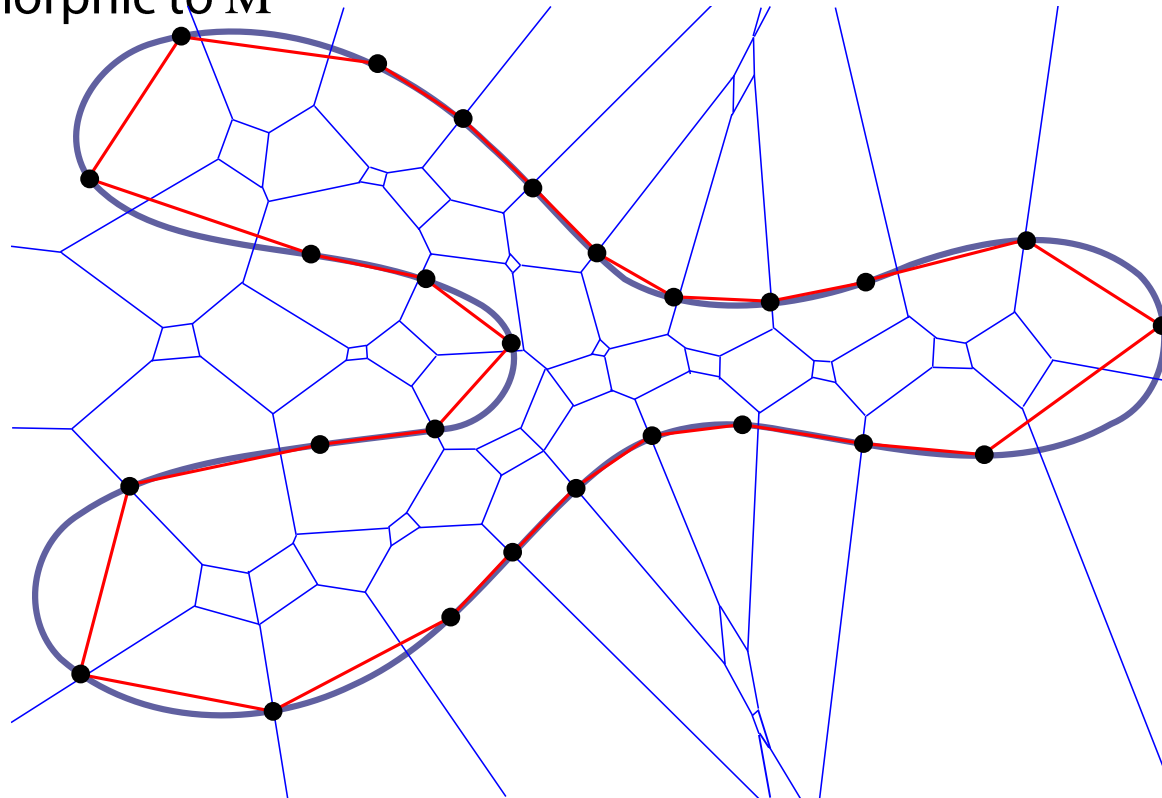


Witness complex

■ Witness complex and the restricted Delaunay complex

[Attali et al., 2007]

There exists a positive ε such that the restricted Delaunay complex for an ε -sample coincides (in the limit) with the witness complex and both are homeomorphic to M



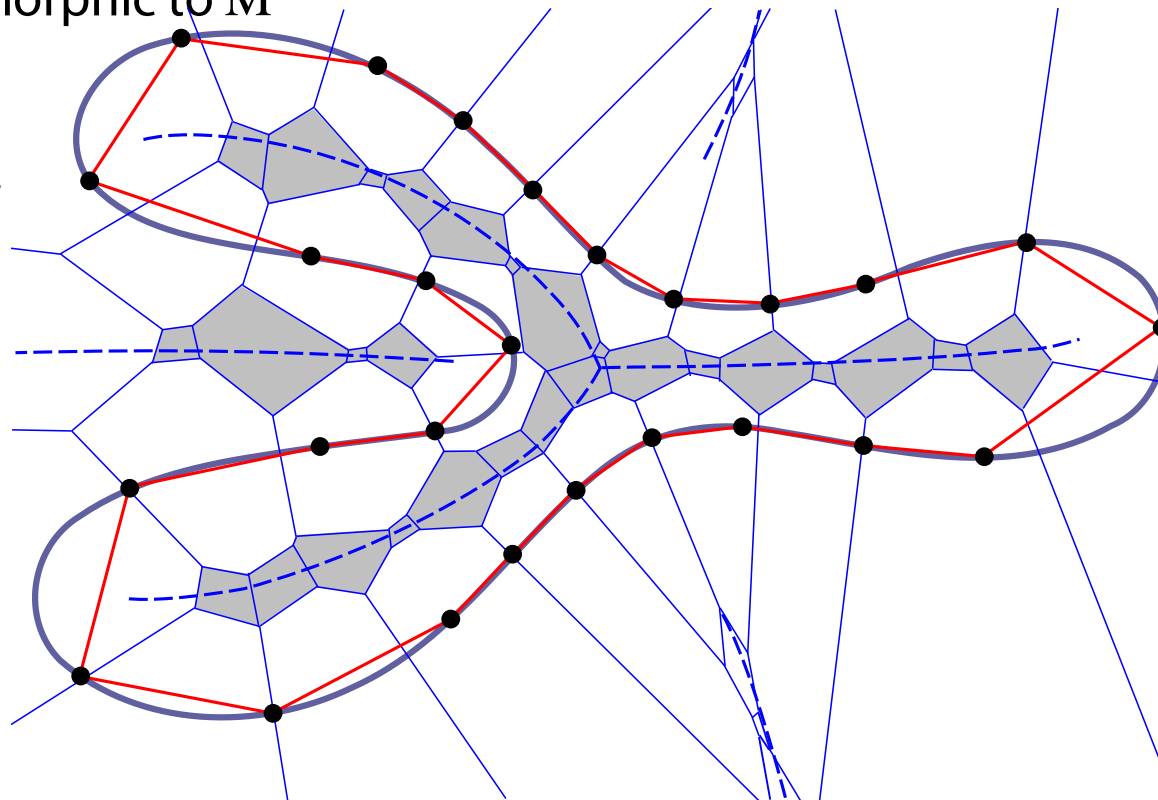
Witness complex

▪ Witness complex and the restricted Delaunay complex

[Attali et al., 2007]

There exists a positive ε such that the restricted Delaunay complex for an ε -sample coincides (in the limit) with the witness complex and both are homeomorphic to M

The second-order cells for the "extra" connections tend to aggregate around the medial axis



Self-Organizing Adaptive Map (SOAM)

■ The algorithm

A set L of *units* (aka *landmarks*), initially containing two units only.

Each unit is associated to a few variables:

- 1) A *position* \mathbf{p} in the ambient space
- 2) A *firing counter* f , which decays exponentially with unit activation
- 3) An *activity radius* r
- 4) A *state*, which changes dynamically during the process

A set of *connections* C , initially empty

Each connection is established between two units and is associated to one variable:

- 1) An *age*

A probability distribution $P(\xi)$, having M as its *support*

Self-Organizing Adaptive Map (SOAM)

■ The algorithm

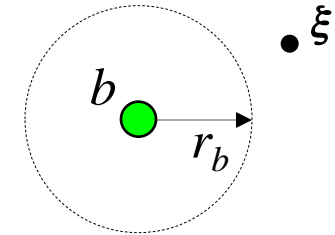
1. Draw a sample ξ from $P(\xi)$
2. Determine the two units b and s whose positions are closest and second-closest to ξ
3. Add the connection (b, s) with $age = 0$ to C , if it is not already present. Otherwise, set its age to 0
4. Unless unit b is in a *stable* state (see below) increase by one the age of all connections involving b .
Remove all connections whose age exceeds a threshold T_{age}
Remove all units that became unconnected, due to this

Self-Organizing Adaptive Map (SOAM)

■ The algorithm

5. If unit b is at least in the *habituated* state and the distance between the input ξ and its position \mathbf{p}_b exceeds its *activity radius* r_b

- create a new unit n
- set its position to \mathbf{x}
- remove the connection (b, s)
- add new connections (b, n) and (n, s)



6. Decrease exponentially the *firing counters* of unit b and of all units connected to it

$$\Delta f_b = (\alpha_h \cdot (F - f_b) - 1) / \tau_f$$

$$\Delta f_{nb} = (\alpha_h \cdot (F - f_{nb}) - 1) / \tau_{f,n}$$

where F is the initial value and the α 's and τ 's are suitable constants

Self-Organizing Adaptive Map (SOAM)

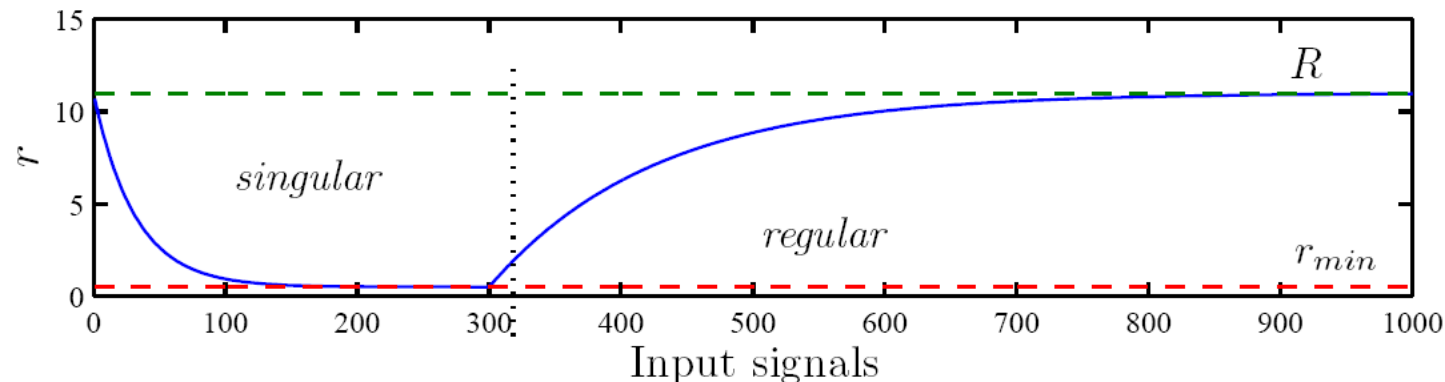
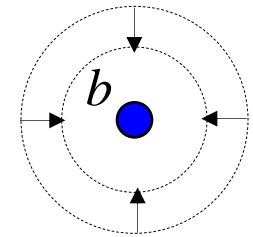
■ The algorithm

7. Update the state of unit b , according to the value of the *firing counter* f_b and the topology of its *neighborhood* of connected units (see below)
8. If unit b is in a *singular* state, decrease exponentially its *activity radius* r_b

$$\Delta r_b = (\alpha_r \cdot (R - r_b) - 1) / \tau_{r, hab}$$

otherwise, if unit b is in a *stable* state increase exponentially r_b

$$\Delta r_b = ((\alpha_r / \tau_{r, dis}) \cdot (R - r_b))$$



Self-Organizing Adaptive Map (SOAM)

■ The algorithm

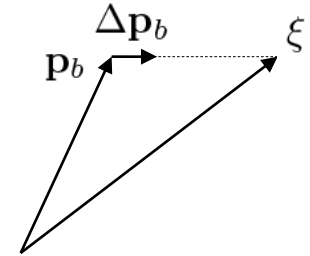
9. Unless unit b is in a *stable* state, adapt its position and those of all connected units

$$\Delta \mathbf{p}_b = \eta_b \cdot f_b \cdot (\xi - \mathbf{p}_b)$$

$$\Delta \mathbf{p}_{nb} = \eta_{nb} \cdot f_{nb} \cdot (\xi - \mathbf{p}_{nb})$$

otherwise, if unit b is *stable*, adapt only the position of b itself

$$\Delta \mathbf{p}_b = \eta_{stable} \cdot f_b \cdot (\xi - \mathbf{p}_b)$$

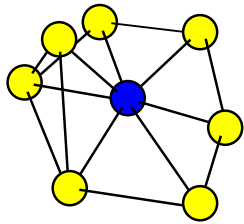


10. Unless some termination criterion has been met, return to step 1.

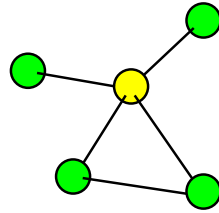
Self-Organizing Adaptive Map (SOAM)

■ Unit states and neighborhood topology

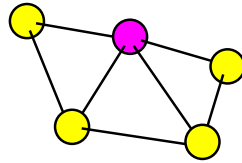
For *surface* reconstruction



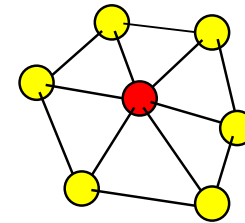
singular
the configuration of
connected
neighboring units
exceeds a disk



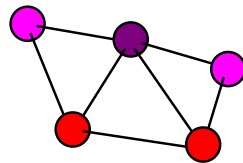
connected
the neighboring units
are habituated



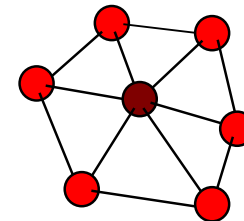
half-disk
formed by connected
neighboring units



disk
formed by connected
neighboring units



boundary
an *half-disk*
formed by regular
neighboring units

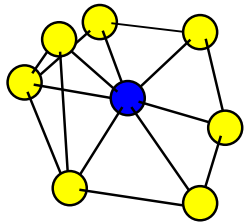


patch
a *disk*
formed by regular
neighboring units

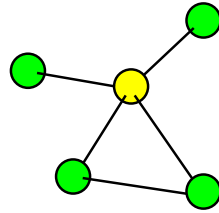
Self-Organizing Adaptive Map (SOAM)

Unit states and neighborhood topology

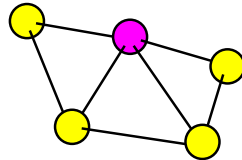
For *surface* reconstruction



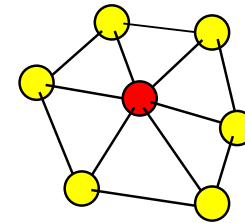
singular
the configuration of
connected
neighboring units
exceeds a disk



connected
the neighboring units
are habituated

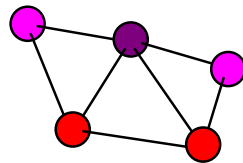


half-disk
formed by connected
neighboring units

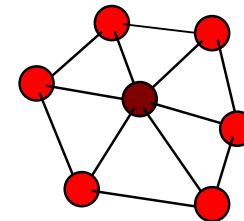


disk
formed by connected
neighboring units

These states
are deemed
stable



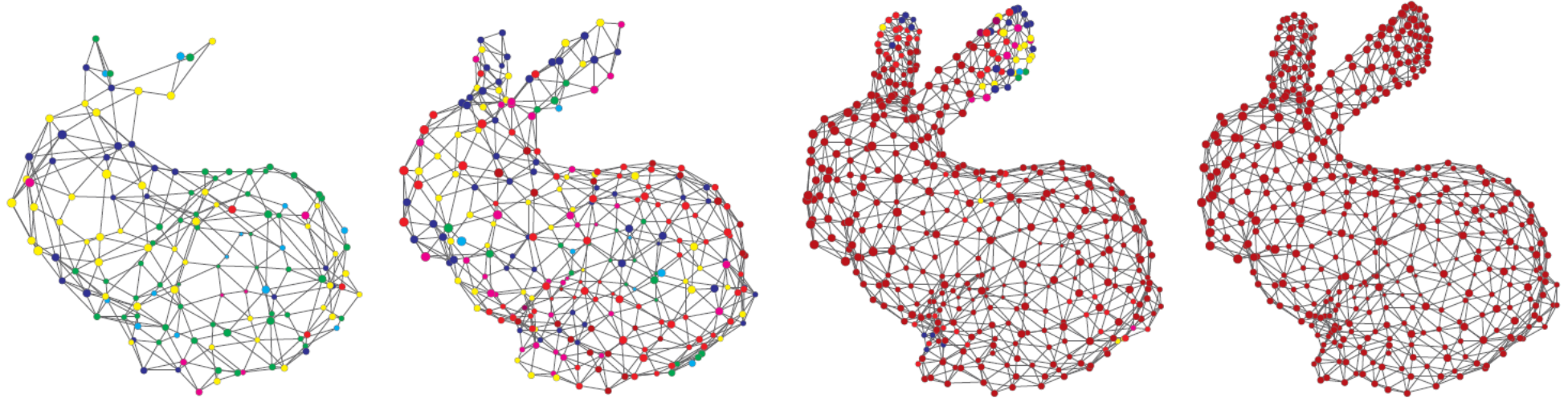
boundary
an half-disk
formed by regular
neighboring units



patch
a disk
formed by regular
neighboring units

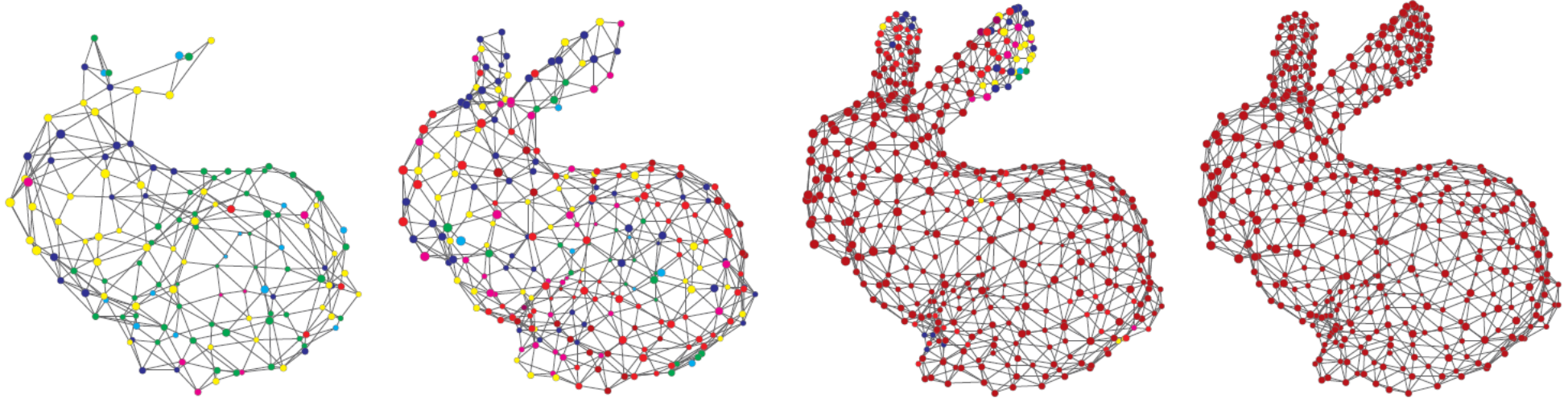
Self-Organizing Adaptive Map (SOAM)

■ SOAM adaptation process

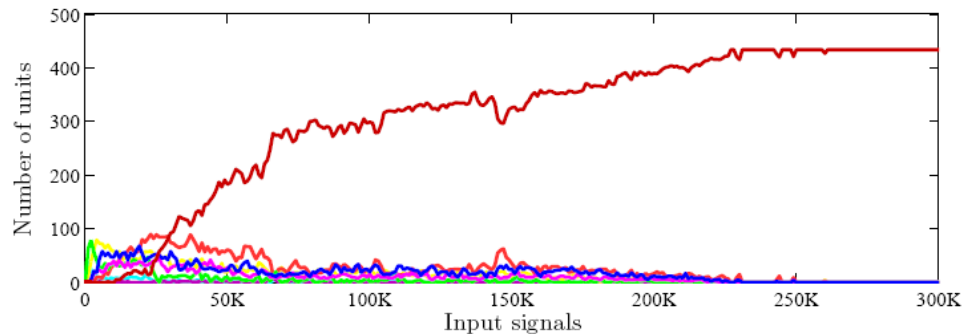


Self-Organizing Adaptive Map (SOAM)

SOAM adaptation process



How the number of units varies with time (i.e. input signals)

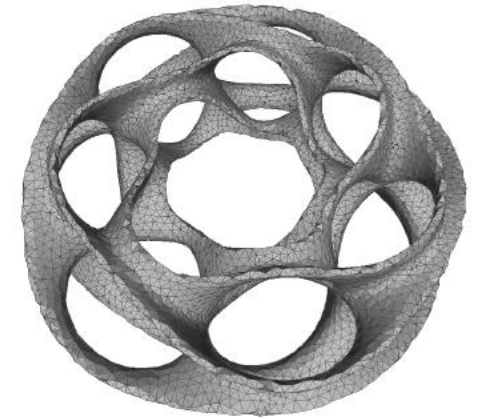
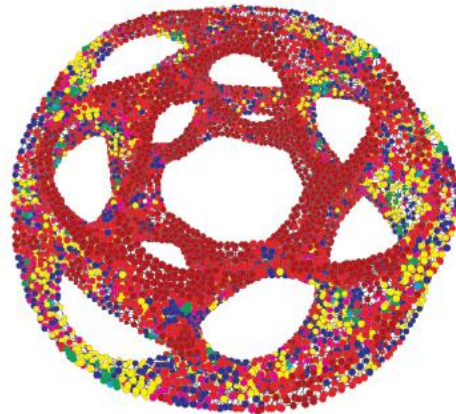


Each line describes the number of units in the corresponding state/color

Self-Organizing Adaptive Map (SOAM)

- **SOAM adaptation process**

Another example, a closed surface with genus 22



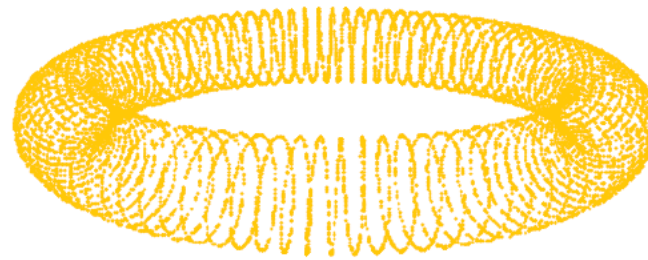
*The same network
interpreted as a mesh*

Self-Organizing Adaptive Map (SOAM)

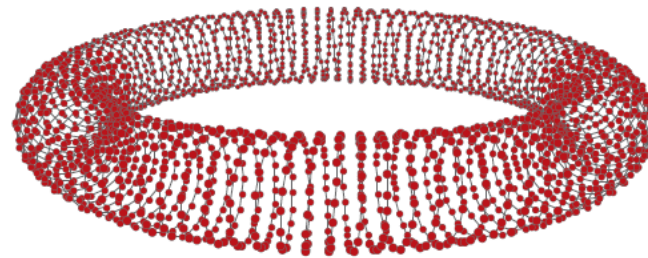
■ SOAM adaptation process

Either a curve or a surface from the same input

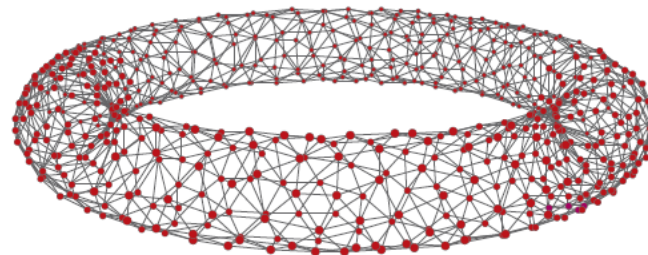
The dimension of the manifold to be reconstructed (i.e. either 1 or 2) is the main parameter of the algorithm



This is the input dataset



The SOAM reconstructs a curve in 3D

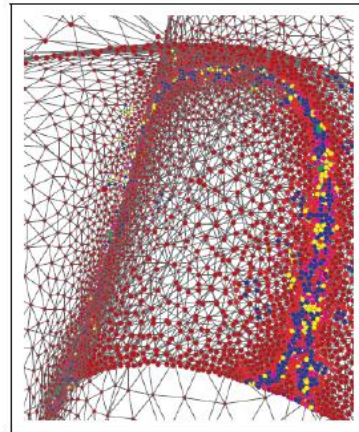
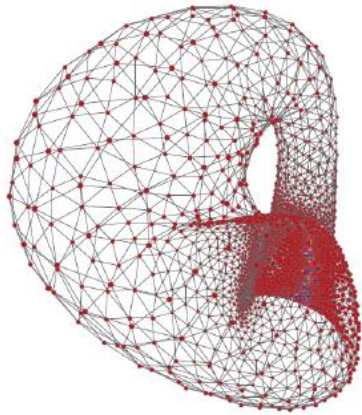


The SOAM reconstructs a surface in 3D

Self-Organizing Adaptive Map (SOAM)

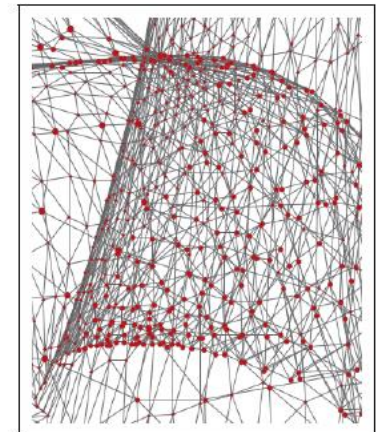
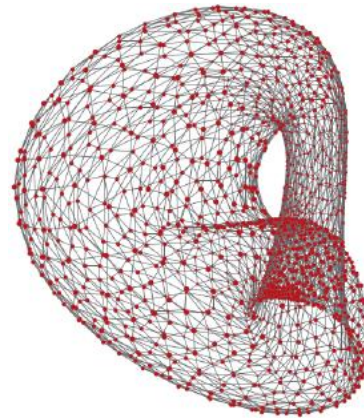
■ SOAM adaptation process

Higher dimensions (i.e. beyond 3D)



In 3D the Klein bottle is not a manifold, as it must self-intersect: the SOAM cannot converge

In 4D (and beyond) the Klein bottle is a manifold and the SOAM converges



Self-Organizing Adaptive Map (SOAM)

- **Pre-print**

See <http://arxiv.org/abs/0812.2969>