

TDA-fall 2025

Monodromy



## Last time: Directional Transforms

Take  $A \subseteq \mathbb{R}^d$ , + compute topological  
signature (ECC, PD, Reeb graph) for  
sublevelset filtration for every direction  
 $w \in S^{d-1}$ .

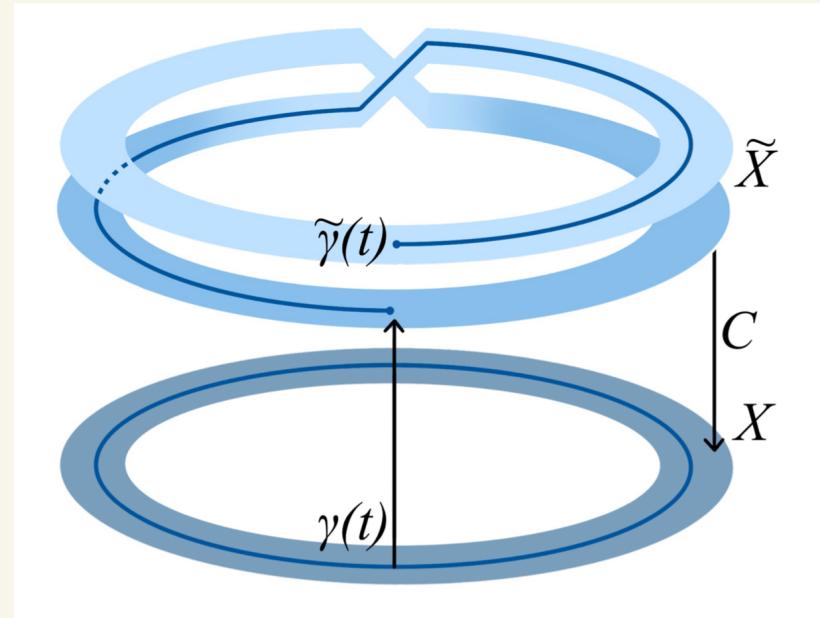
Big result/surprise:

**Monodromy**: effect where loops in a base space don't lift to loops in a covering space

Consider space  $X$ , loop  $\gamma$ , covering  $\tilde{X}$ , and a lift.

If  $\tilde{\gamma}(0) \neq \tilde{\gamma}(2\pi)$ , then  $\gamma$  exhibits monodromy.

(Usually studied for polynomials on points in  $\mathbb{C}^*$ )

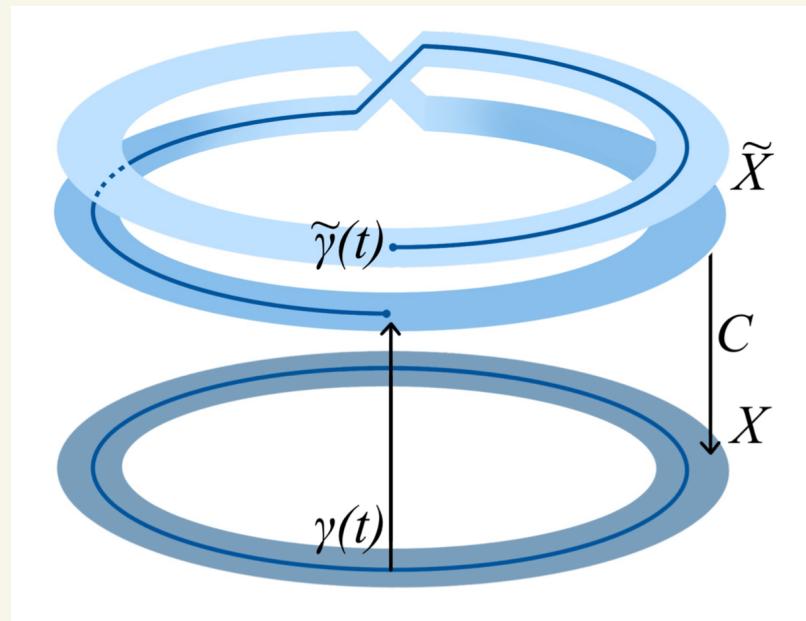


We say  $\gamma$  has monodromy of order  $k$  if  $k$  repetitions of the covering loop returns to the starting point, &  $k$  is minimal such value!

$$\tilde{\gamma}(0) = \tilde{\gamma}^k(k2\pi)$$

$$= \tilde{\gamma}_0 \cdots \circ \tilde{\gamma}$$

$k$  times



Here:  $k=2$

# Monodromy & directional transforms

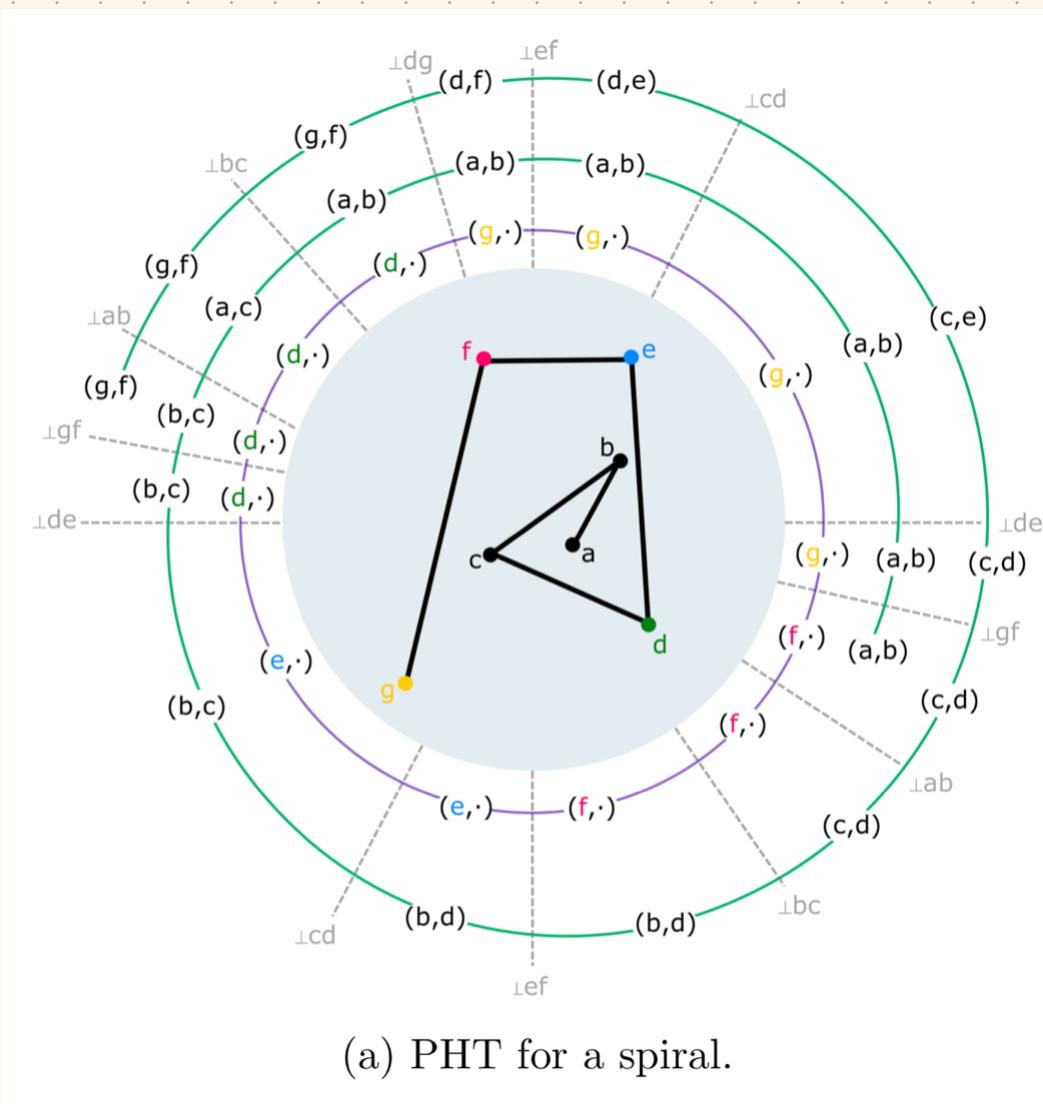
Recently

Ayer et al 2024

studied

monodromy

for the directional transform:



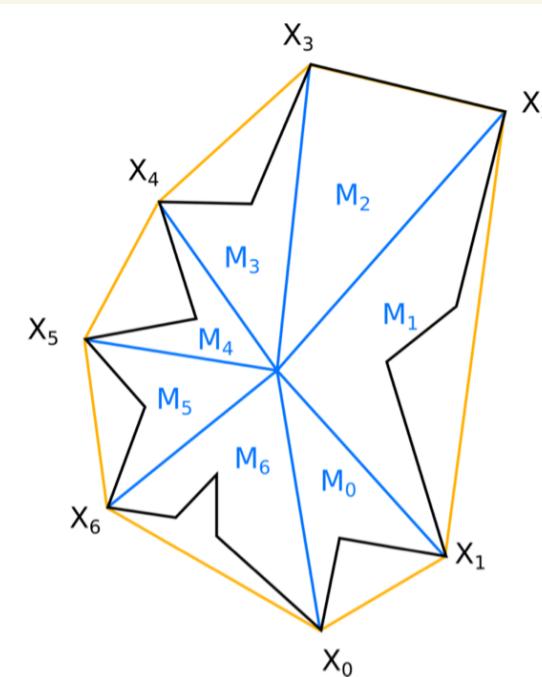
## Star-Shaped objects in $\mathbb{R}^2$

A shape  $M$  is star-shaped if  $\exists$  point  $c \in M$  such that for any  $x \in M$ , the line is contained inside of  $M$ .

[Arya et al] prove that, if  $A \subseteq \mathbb{R}^2 + A$  star-shaped  $\Rightarrow \text{PHT}(A)$  has no monodromy.

Proof technique: decompose into sectors, & Show

$$\text{PHT}_0(A) \cong \bigoplus_{\substack{\text{sectors} \\ M_i}} \text{PHT}_0(M_i)$$



## Generalizing

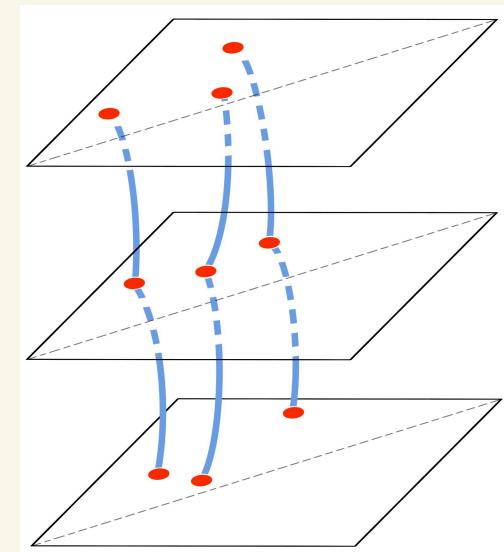
Unfortunately, unclear if reverse holds,  
so not (yet) a complete characterization  
of what monodromy is catching.

In  $\mathbb{R}^d$  for  $d > 3$ , proof does not work:  
there are star-shaped objects where  
 $\text{PH}_0$  cannot be decomposed into sectors.

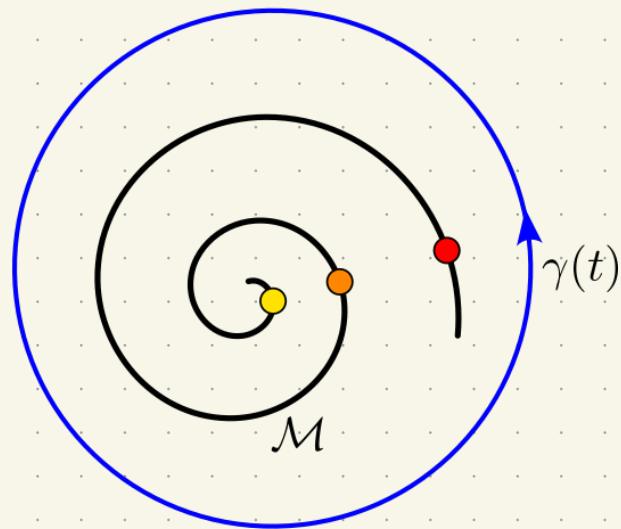
Other direction: how much monodromy can we find?

Vineyard viewpoint:

Monodromy is a permutation  
of points in "z-axis"



Back to that spiral:



A different transform

Onus et al

Let  $d(\cdot, x) : M \rightarrow \mathbb{R}$  be the distance

from any  $x \in \mathbb{R}^d$  to  $M \subset \mathbb{R}^d$

↳ called radial distance fan

Fix a loop  $\gamma : [0, 2\pi] \rightarrow \mathbb{R}^d$

↳ observation loop

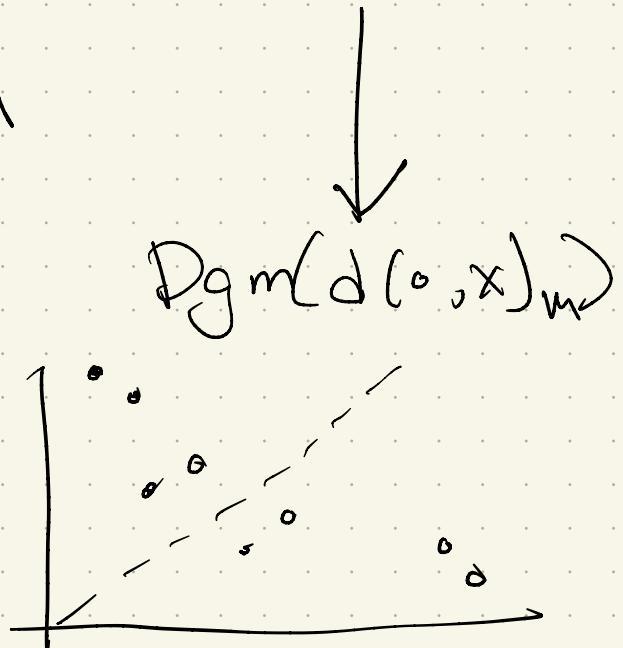
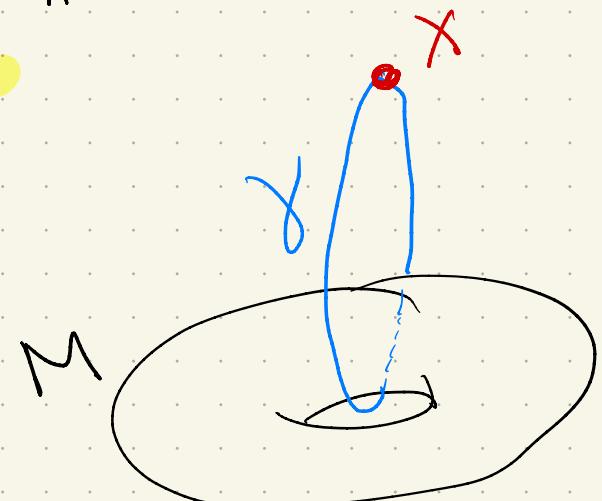
Set  $x = \gamma(t)$  for  $t \in [0, 2\pi]$

↳ family of filtrations  $d(\cdot, \gamma(t))_M$

Closed Vineyard map

$CV_M : S^1 \rightarrow S^1 \times Dgm$

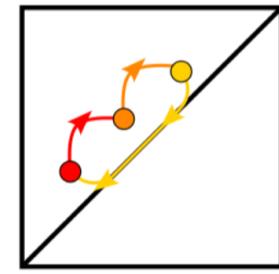
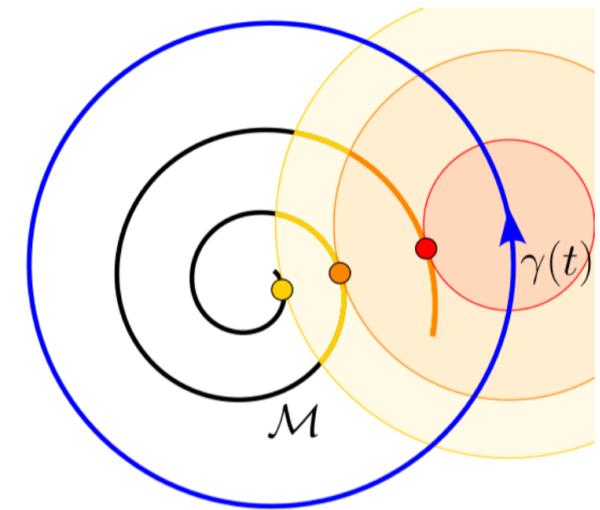
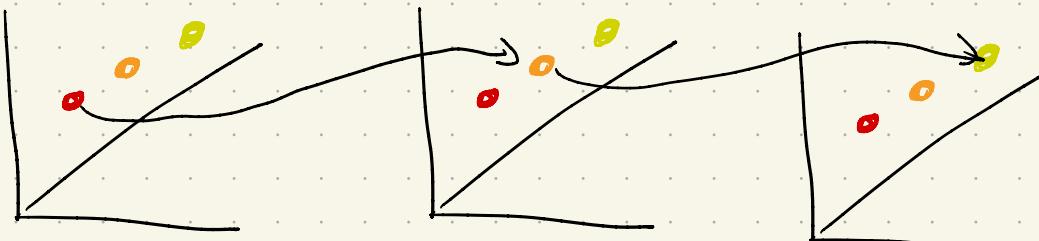
$t \mapsto (t, Dgm(d(\cdot, \gamma(t))_M))$



# Re-doing the spiral

In this example, the vines will still induce a map from  $\text{Dgm}(\text{d}(\cdot, \gamma(t))_M)$  to 'itself' which permutes the points

Vineyard:



$\text{Dgm}_0(\text{d}(\cdot, \gamma(t))_M)$

Here, we have monodromy of order 3.

Arya et al 2024

Can we demonstrate monodromy in  
higher dimensions?  
Which objects have it, & how much?

Theorem

C., Fillmore, Stephenson, Wintzreeden

Monodromy of any order  $k$  can be  
created in the  $\ell$ -vineyard of the  
radical transform of a manifold  $M$   
embedded in  $\mathbb{R}^{l+1}$

## Idea for how to prove

We know the radial vineyard can reflects parts of the topology of input space  $M$ .

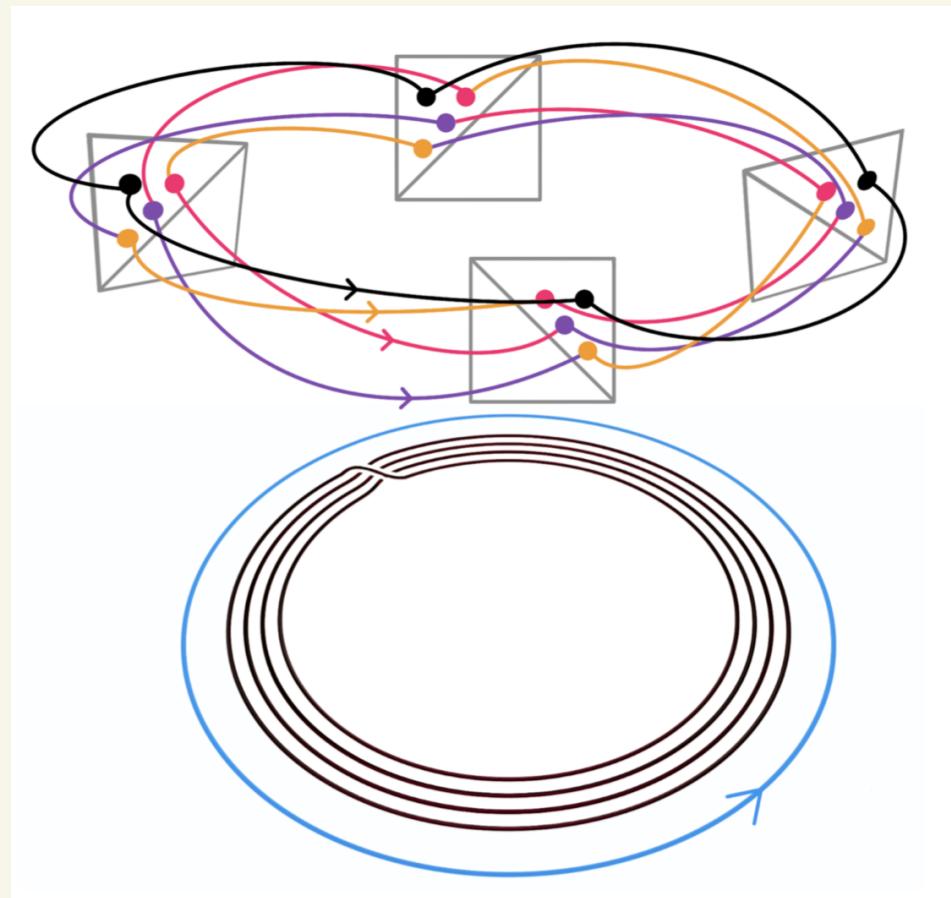
If  $M$  were a braid, with period  $2\pi k$ , we could get order  $k$  monodromy.

Braids have previously connected to monodromy.

Cohen & Scau 1997,

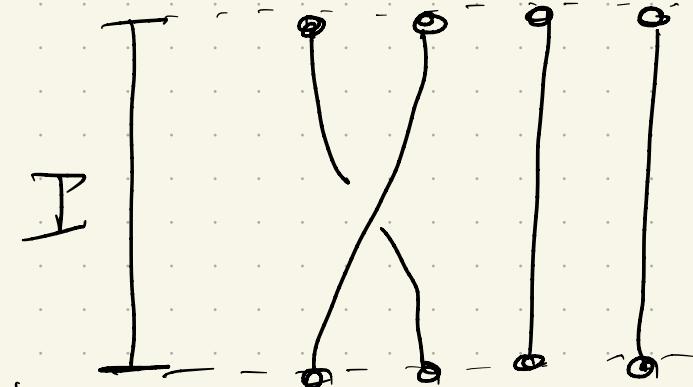
Cogolludo-Agustin 2011

Salter 2023, Salter 2024



# Brands

A braid on  $m$  strands  
is the equivalence class of  
the disjoint union of  $m$   
intervals  $B_i : \mathbb{I} \rightarrow D^2 \times \mathbb{I}$ , monotonically  
increasing wrt  $\mathbb{I}$ , such that end points are  
a permutation of start points, under  
ambient braid isotopy.



Variant of  
Reidemeister  
moves



Figure 4: The Reidemeister moves. Left to right: Type I, Type II, Type III.

## Composing braids:

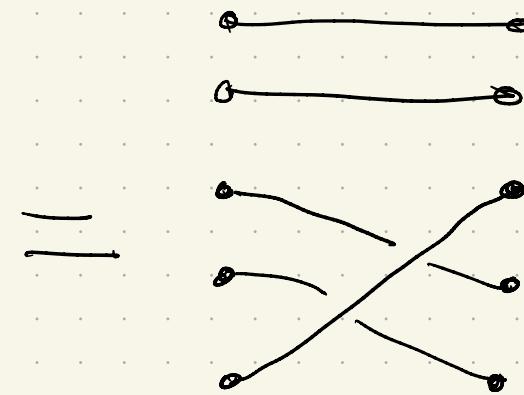
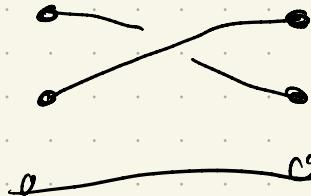
$B_1$



$B_2$

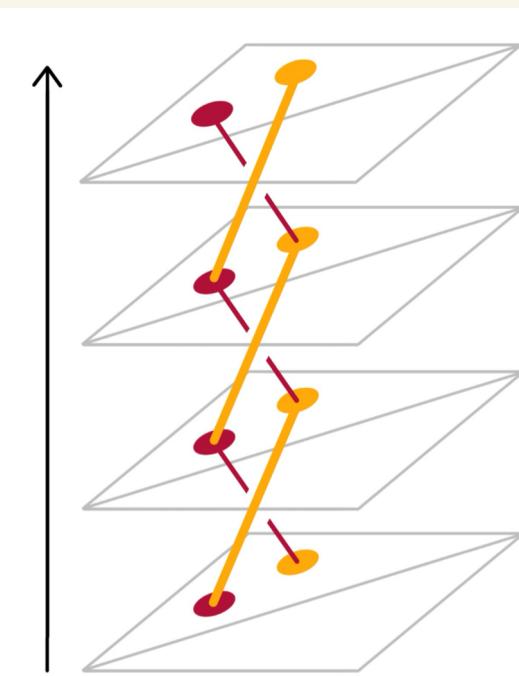


+



If we look at vineyards,  
can see some of the  
same patterns

Let's can ask if they  
contain braids.

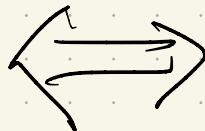
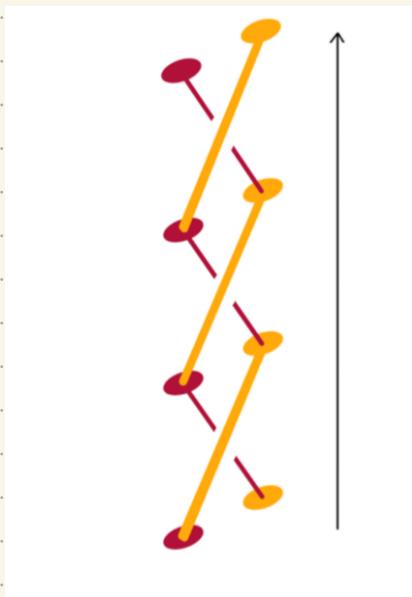


# Braids & knots

If we identify endpoints of a braid & map canonically to torus in  $\mathbb{R}^3$ , we get a closed braid.

Theorem [Alexander 1923]

Every knot or link is equivalent to a closed braid.

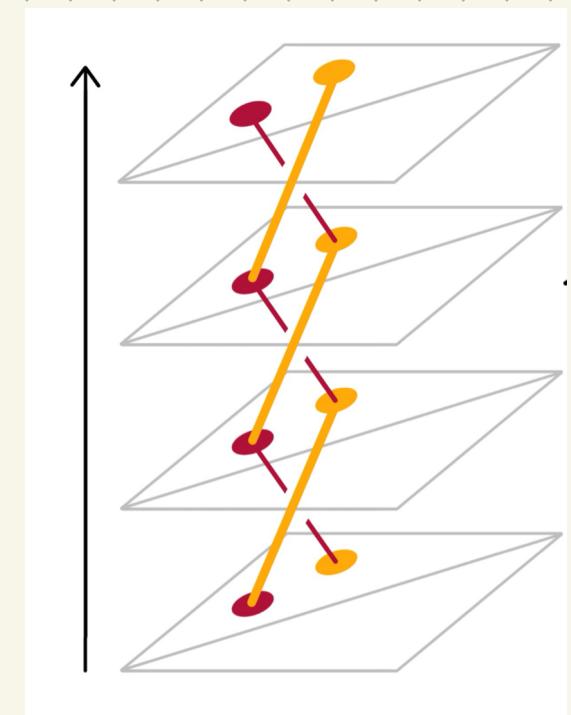
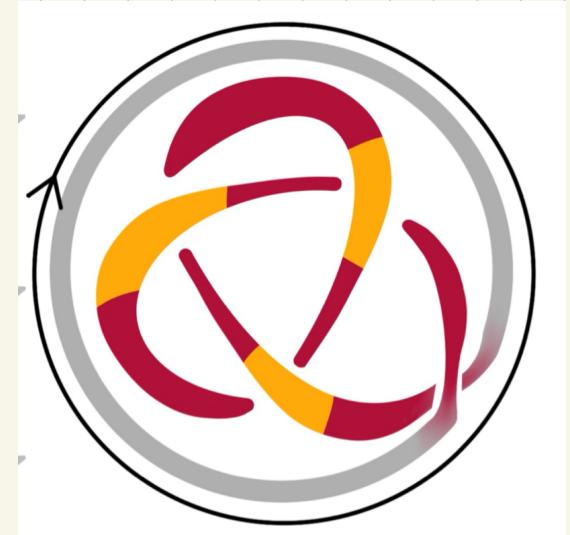


**Question:** Can vineyards "braid"?

**Theorem**

C.-Fillmore-Stephenson-Wintrebecken

Given any braid  $B$ , there exists a manifold  $M \subset \mathbb{R}^d$  & a closed curve  $\gamma \subset M$  such that identifying the ends of the vineyard of  $d(\cdot, \gamma(t))_M$  will yield a braid  $B'$  which is equivalent to  $B$  after removing spurious components.



## Construction

Note that both  $M$  and  $\gamma$  must be carefully constructed to work together!

## Overview:

Start with a closed braid  $B \subset R^2$ , with  $k$  components and  $s$  strands.



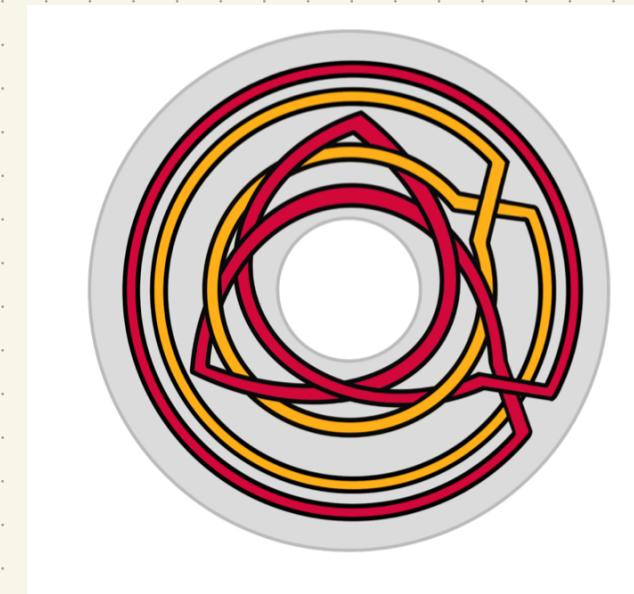
[Note: Not showing the braid here, but  $k=2$  and  $s=3$ ]

We'll convert this to a manifold in  $R^3$

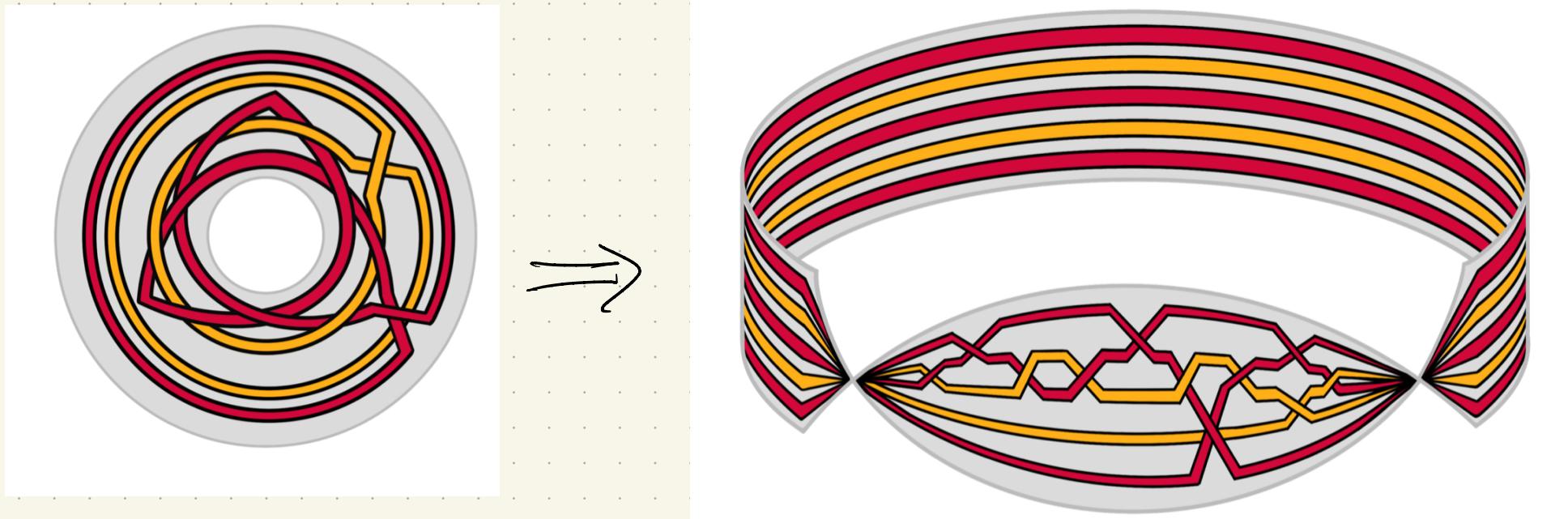
① Redraw in a small neighborhood of an annulus, where strands follow fixed radii

② Then, introduce an extra "twist" per component & wrap loop around outside of annulus

⇒ adds  $O(sk)$  crossings  
and gives  $n = s+k$   
strands total



- ③ Deform B so all crossings occur in fraction of annulus, at regular intervals
- ④ Twist annulus in 3d, so non-crossing part is orthogonal. (Note: no new crossings!)
- ⑤ Set observation loop to follow annulus at fixed distance + near "center"

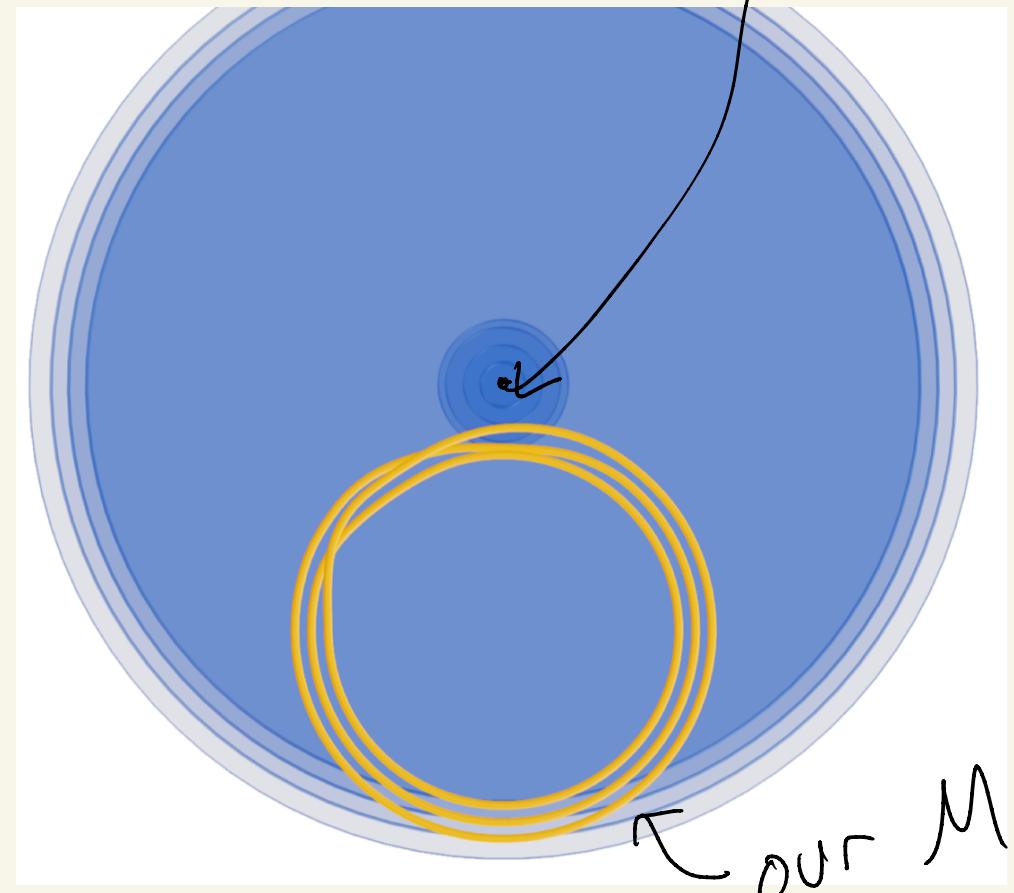


## The vineyard

Several technical lemmas relying on Morse theory + angles show that births in  $H_0$  all occur before deaths:

point on  $\gamma$

This means we can use embedding to control vineyard diagrams.



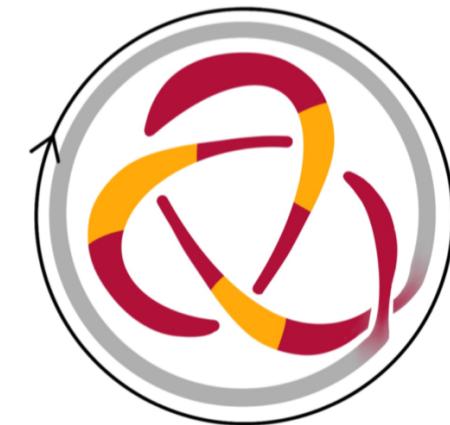
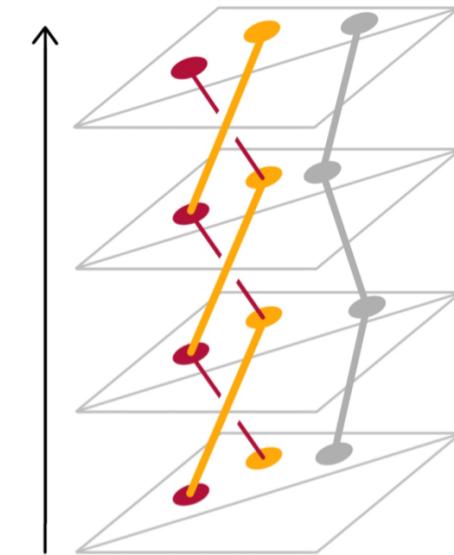
Back to that extra loop

Elder rule in persistence

⇒ first birth & last  
death are paired.

We added an outer strand  
to account for this.

Result: For each component,  
will be an unlinked  
strand, which will be  
an extra circle in  
vneward.

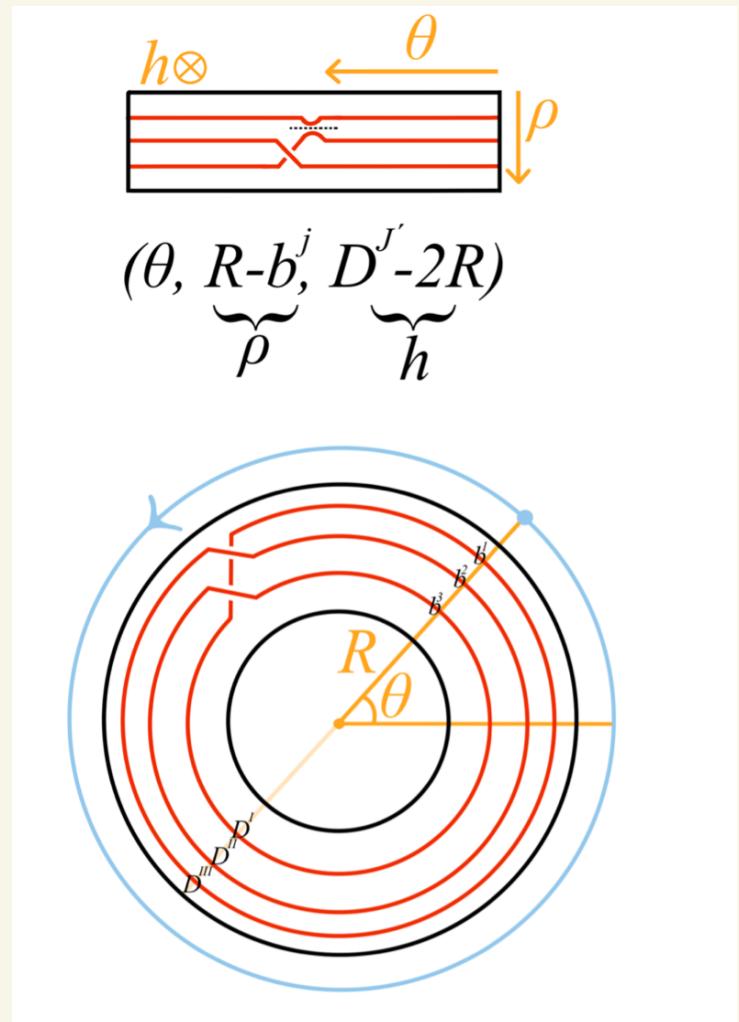


# Parameterizing the vineyard

Our vines have annular coordinates:  
 $\theta, h \& P$

Away from crossings,  
vines have distinct  $P$   
so  $h$  doesn't matter.

To get correct over/under  
crossings in vineyard,  
we need to play with  
the geometry of our embedding a bit...

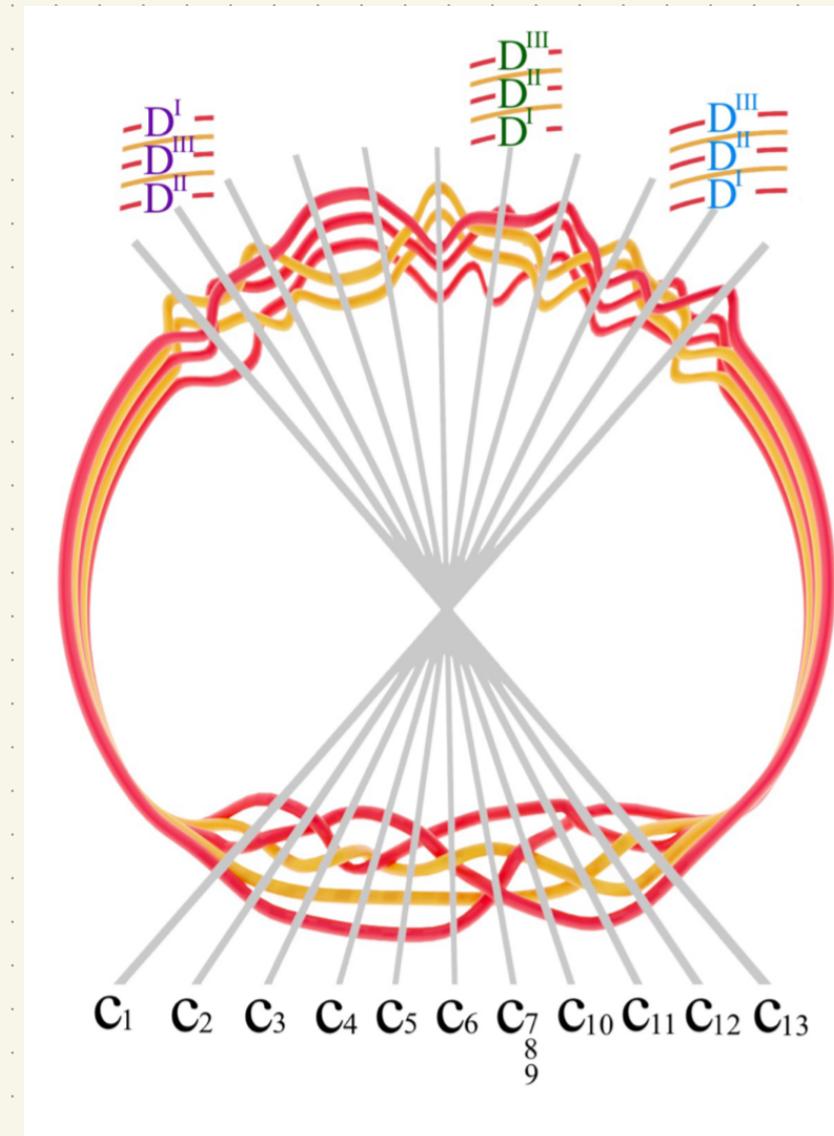


# Pushing or Pulling

Our construction ensured that crossings are evenly spaced, & all deaths occur at opposite points.

Therefore, we perturb deaths to make sure Vineyard has correct crossings.

(Note: hiding some intense calculations here.)



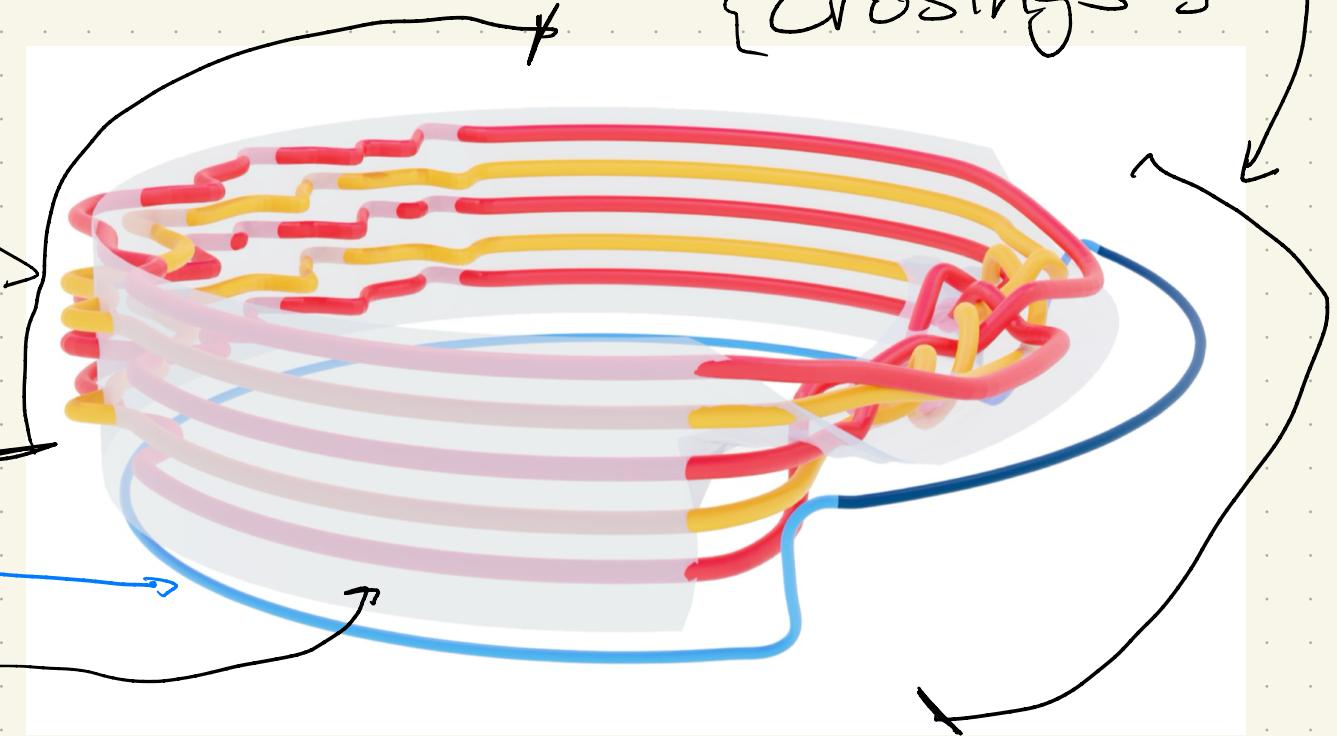
Final result: braids

{ all crossings }

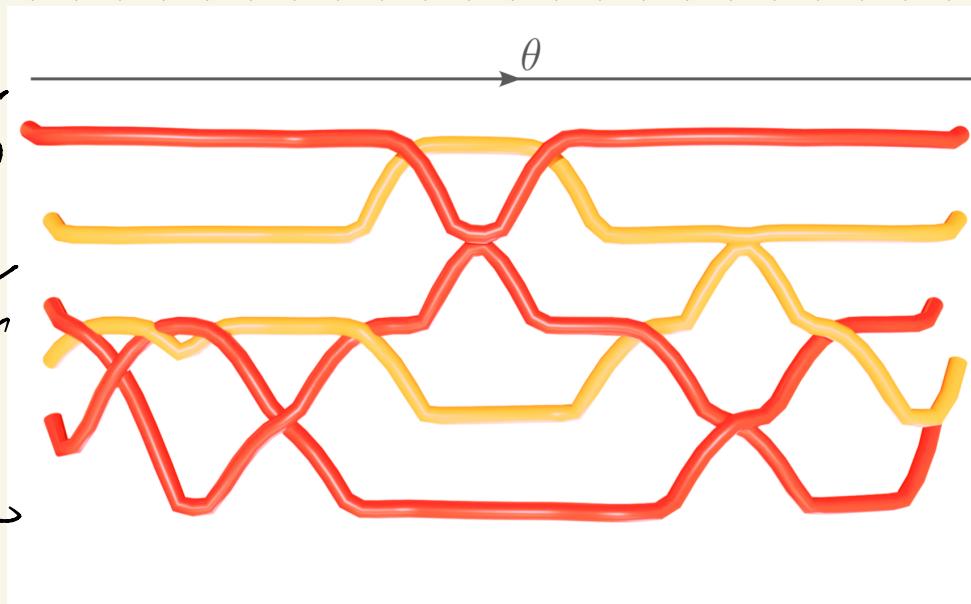
Pushed &  
pulled  
deaths

observations  
loop

base annulus



unlinked  
outer  
loops  
braid?



Vineyard of dark  
blue portion of  
the observation  
loop

## Higher Dimensions

To have this calculation work for higher dimensions, take  $(l+1)$ -dimensional  $\alpha$ -offset of braid  $B \times O$  in  $R^3 \oplus R^{l-1} \subset R^d$ .

Given this manifold construction  
(+ same base loop), vines in our  
 $l$ -vineyard will be  $\alpha$ -close to the  
 $O$ -vineyard.

So why study vineyards?

Recognizing knots is not easy

↳ but classifying vineyards might involve knots

That said, classifying knots is very well studied!

Other (new!)  
approaches  
ignore the  
knots.

vy1 [math.AT] 28 Oct 2025

## Through the Grapevine: Vineyard Distance as a Measure of Topological Dissimilarity

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October 29, 2025

### Abstract

We introduce a new measure of distance between datasets, based on vineyards from topological data analysis, which we call the vineyard distance. Vineyard distance measures the extent of topological change along an interpolation from one dataset to another, either along a pre-computed trajectory or via a straight-line homotopy. We demonstrate through theoretical results and experiments that vineyard distance is less sensitive than  $L^p$  distance (which considers every single data value), but more sensitive than Wasserstein distance between persistence diagrams (which accounts only for shape and not location). This allows vineyard distance to reveal distinctions that the other two distance measures cannot. In our paper, we establish theoretical results for vineyard distance including as upper and lower bounds. We then demonstrate the usefulness of vineyard distance on real-world data through applications to geospatial data and to neural network training dynamics.

- Decompose into vines
- Project vines into  $xy$  plane
- Measure length there

