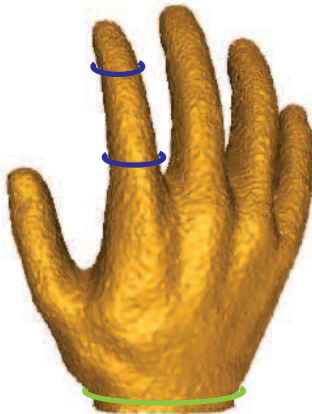


# Computing Optimal Homotopies

Erin Wolf Chambers  
Saint Louis University  
erin.chambers@slu.edu

# Motivation: Measuring Similarity Between Curves

How can we tell when two cycles or curves are similar to each other?



Similarity measures have many potential applications:

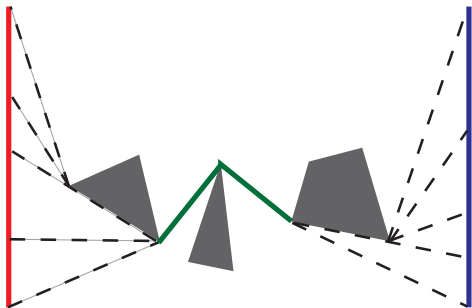
- Analyzing GIS data
- Map analysis and simplification
- Handwriting recognition
- Computing “good” morphings between curves
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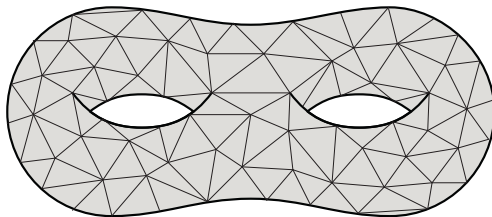
There are many different ways to check similarity. Most focus on either the geometry or the topology of the curve and the ambient space.

I'll consider two different settings (at least for the most part). First setting:

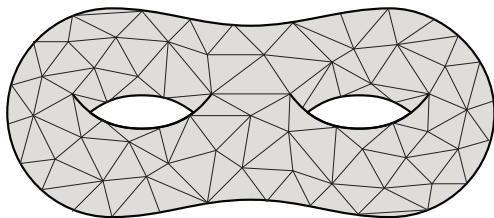


The plane, sometimes minus a set of (polygonal) obstacles.

Second setting: A combinatorial or piecewise linear orientable surface.



Second setting: A combinatorial or piecewise linear orientable surface.



Any such space is homeomorphic to a sphere with a number of handles attached; we call this number the *genus* of the surface.

# Hausdorff distance

Hausdorff distance is one way to measure distance: consider any point on one curve and find the closest point on the other curve. Take the largest of all these distances.

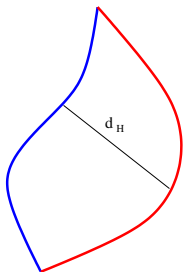


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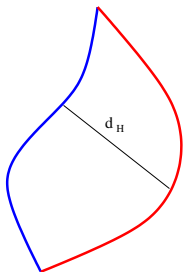


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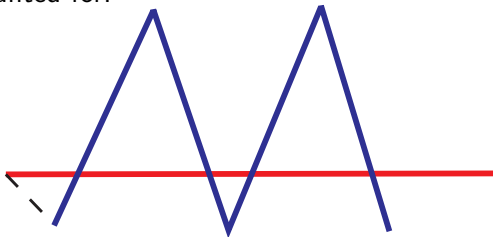
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# Fréchet Distance

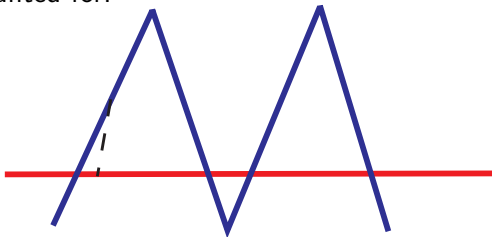
With Fréchet distance (or dog leash distance), the flow of the curve is accounted for.



Imagine a man walking along one curve and a dog along the other, with a leash always connecting them, and minimize the length of the longest leash.

# Fréchet Distance

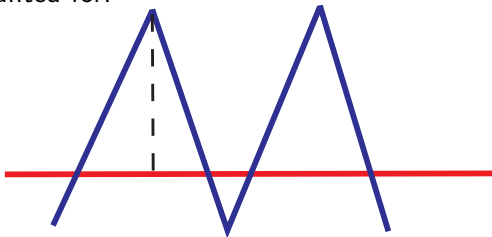
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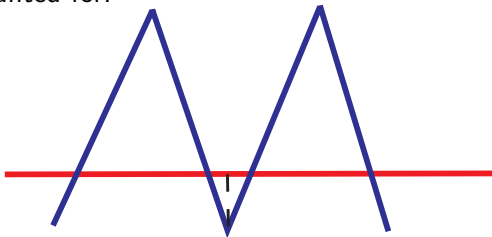
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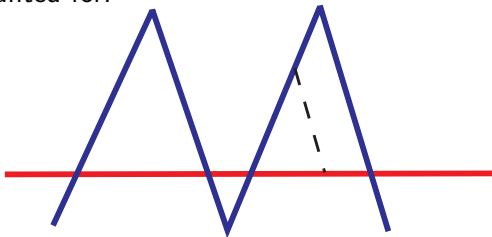
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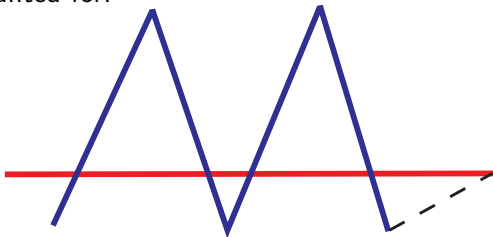
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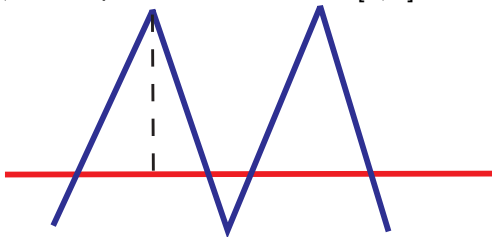


# Fréchet Distance

More formally, given two curves  $\gamma_1$  and  $\gamma_2$ , the Fréchet distance is:

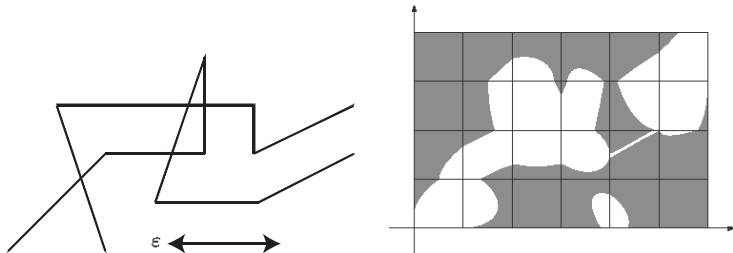
$$F(A, B) = \inf_{\alpha, \beta} \max_{t \in [0,1]} \{d(\gamma_1(\alpha(t)), \gamma_2(\beta(t)))\}$$

where  $\alpha$  and  $\beta$  are reparameterizations of  $[0, 1]$ .



# Main tool: Free space diagram

Consider each pair of segments from the two curves, and calculate which portions are within  $\epsilon$  of each other.



We build the *free space diagram* by forming the  $n$  by  $m$  grid, and determine if there is a matching that keeps the leash  $\leq \epsilon$  by searching in this grid.

# Computing the Fréchet distance

Alt and Godau gave the first algorithm to compute this for piecewise linear curves in the Euclidean space: for a fixed  $\epsilon$ , the running time is  $O(mn \log mn)$ .

Parametric search techniques can then be applied to find the best such  $\epsilon$ ; this adds an extra  $O(\log n)$  to the running time.

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Since the initial algorithm, it has been studied extensively: applications, approximations, improved algorithms for restricted classes of curves, and lower bounds are just a few of the many results.

# Fréchet distance in higher dimensions

In addition, Fréchet distance has also been considered in higher dimensions:

- It is NP-Hard to compute the Fréchet distance between two surfaces [Godau 1998].
- Even NP-Hard to compute between terrains or polygons with holes [Buchin-Buchin-Schulz 2010].
- Still NP hard even for surfaces traced by curves [Buchin-Ophelders-Speckmann 2015].
- There is a  $(1 + \epsilon)$ -approximation algorithm for computing Fréchet distance between genus zero surfaces, where the running time is bounded if the input surfaces are “nice” [Nayyeri Xu 2016].

# Geodesic Fréchet Distance

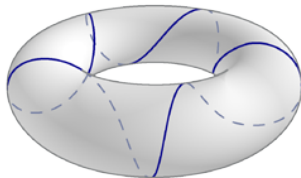
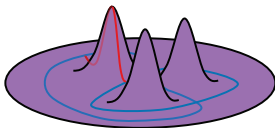
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A geodesic is a path that avoids any obstacles and cannot be locally shortened by perturbations.

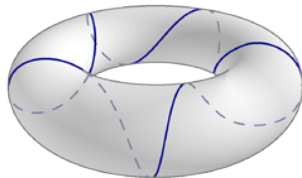
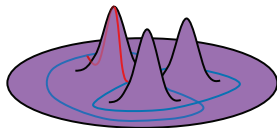


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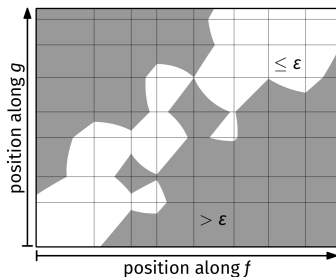
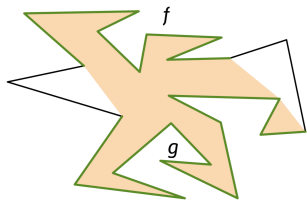
In geodesic Fréchet distance, the leash is required to be a geodesic in the ambient space.



# Geodesic Fréchet Distance

Algorithms are known in some limited settings, such as convex polytopes [Maheshwari and Yi 2005] and simple polygons [Cook Wenk 2008].

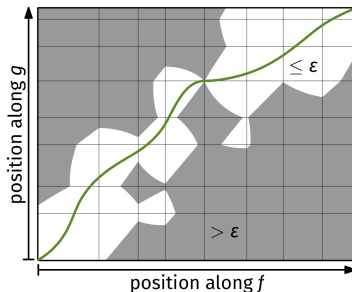
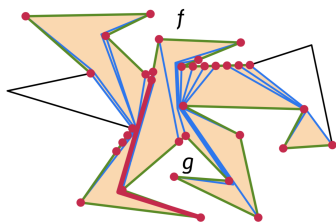
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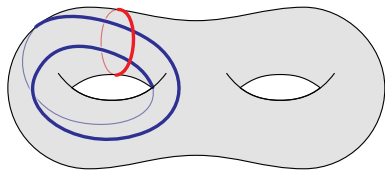
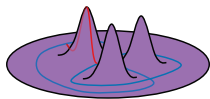


# Adding topology to our measures

So far, all the measures considered have been fairly geometric in motivation: based primarily on actual distances between the points.

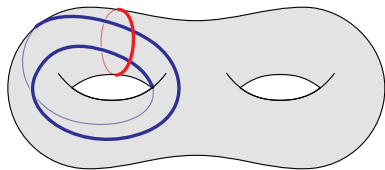
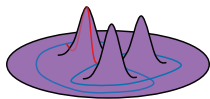
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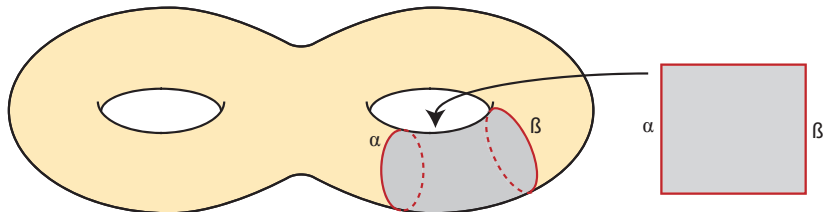


The definition of geodesic Fréchet distance will directly generalize to surfaces, but does not force leashes to move continuously or to be shortest paths.

# Homotopy

## Definition

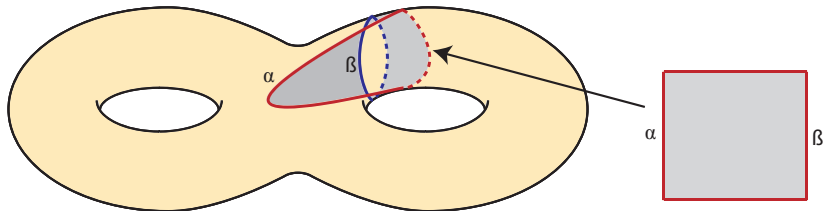
A **homotopy** is a continuous deformation of one path to another. More formally, a homotopy between two curves  $\alpha$  and  $\beta$  on a surface  $M$  is a continuous function  $H : [0, 1] \times [0, 1] \rightarrow M$  such that  $H(\cdot, 0) = \alpha(\cdot)$  and  $H(\cdot, 1) = \beta(\cdot)$ .



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# Testing if two curves are homotopic

Testing if two curves are homotopic has been studied.

- Cabello et al (2004) give an algorithm to test if two paths in the plane minus a set of obstacles are homotopic in  $O(n^{3/2} \log n)$  time; there is also an output sensitive algorithm that takes  $O(\log^2 n)$  time per output vertex [Bespamyatnikh 2003].
- Given a graph cellularly embedded on a surface and two closed walks on that graph, there is an  $O(n)$  time algorithm to decide if the two walks are homotopic [Dey and Guha 1999, Lazarus and Rivaud 2011, Erickson and Whittlesey 2012].



# Combinatorially optimal homotopies

There is work [Chang-Erickson 2016] on finding the “best” homotopy, as well; usually, this involves minimizing number of simplifications moves to untangle a curve.



**Figure 1.1.** Homotopy moves  $1 \rightarrow 0$ ,  $2 \rightarrow 0$ , and  $3 \rightarrow 3$ .

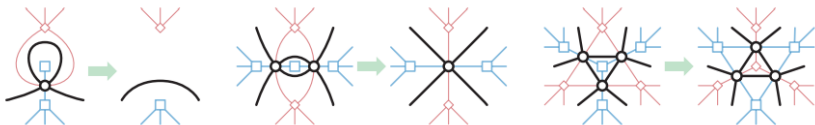
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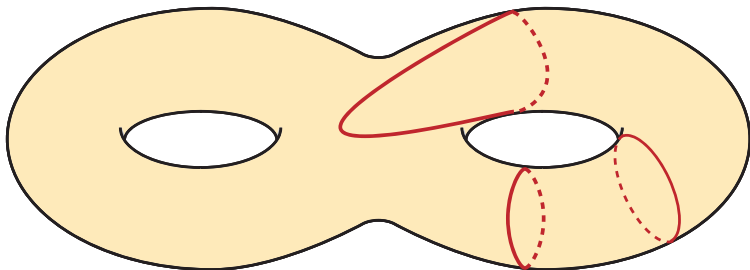
In the plane, they prove this is  $\Theta(n^{3/2})$ .



This connects to older results [Steinitz 1916, Francis 1969, Truemper 1989, Feo and Provan 1993, Noble and Welsh 2000], and electrical moves on the medial graph of the input planar graphs.

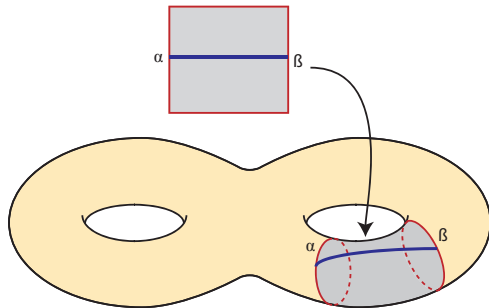
# Beyond testing homotopy

However, in many applications we'd like to include more of a notion of the geometry of the underlying space, as well.



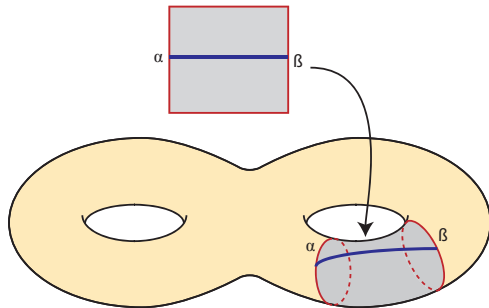
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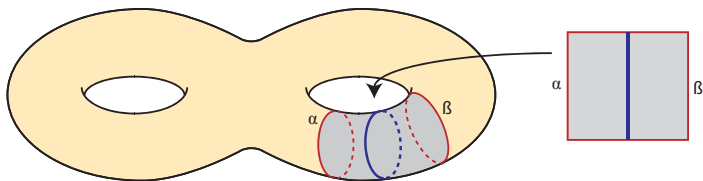


(We could just have really called this the *width* of the homotopy.)

# Height of a homotopy

The height of a homotopy is an orthogonal definition to homotopic Fréchet distance:

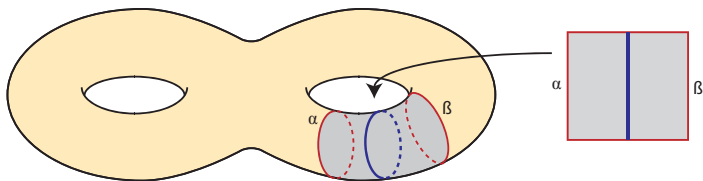
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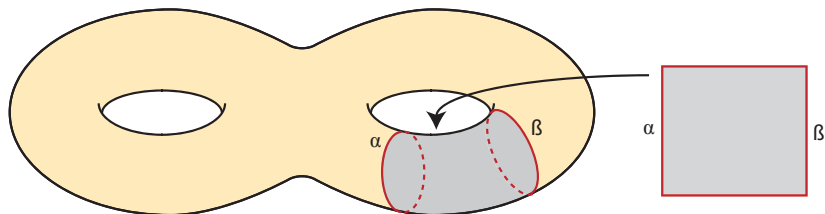
$$d_{HH}(\gamma_1, \gamma_2) = \inf_{\text{homotopies } H} \{ \sup \{ |H(s, \cdot)| \mid s \in [0, 1] \} \}$$



This is also sometimes called an *L-homotopy* [Frosini 1999, G. Chambers-Liokumovich 2014] in the Riemannian setting, or a *B-northward migration* in the combinatorial setting [Brightwell-Winkler 2009].

# Homotopy area

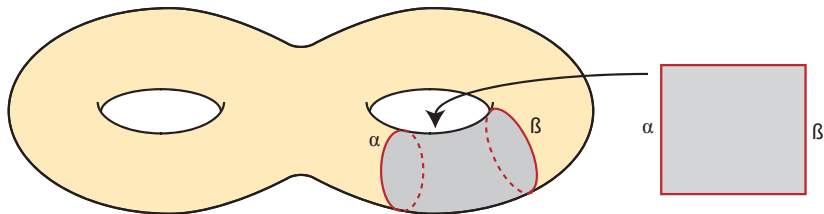
Instead of focusing on the length or width, we can also examine the total area swept by a homotopy or homology.





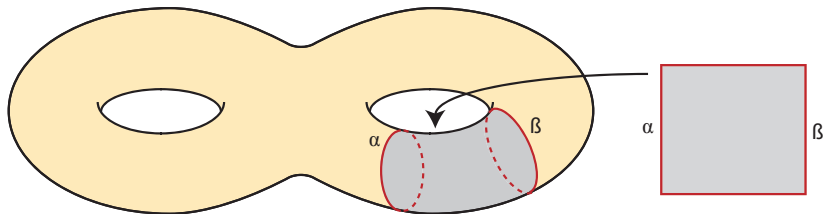
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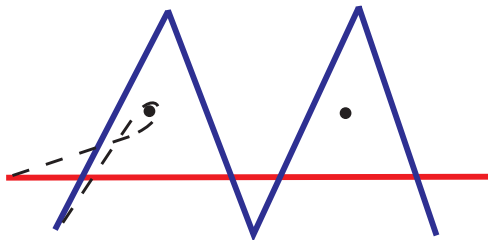


However, homotopy can be notoriously hard to compute! I'll discuss the trade-offs, both in terms of what the homotopy is measuring and (more importantly) how computable the measures are in various settings.

# Homotopic Fréchet Distance

The homotopic Fréchet distance is the length of the shortest leash we can use for our homotopy. Formally,

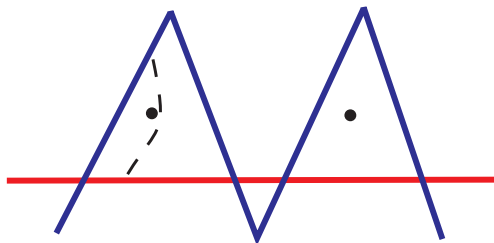
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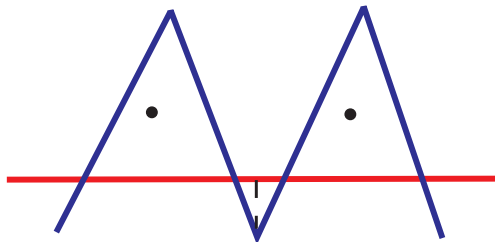
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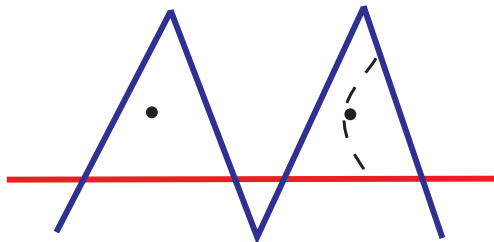
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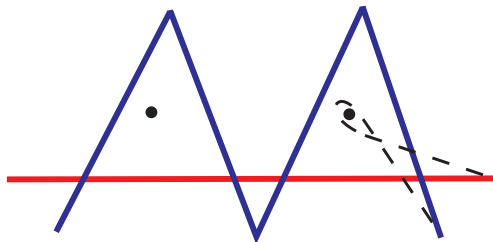
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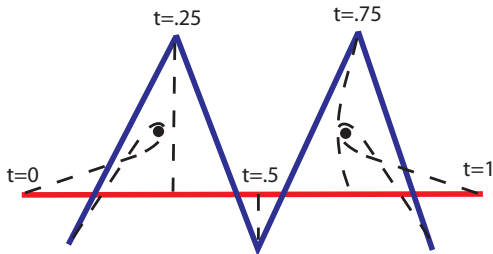
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# Computing the Homotopic Fréchet Distance

There is a polynomial time algorithm algorithm to compute the homotopic Fréchet Distance between two polygonal curves in the plane minus a set of polygonal obstacles [C.-Colin de Verdière-Erickson-Lazard-Lazarus-Thite 2009].

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The algorithm has some similarities to the work of Alt and Godau, but is considerably more complex since there are an infinite number of homotopy classes to consider.

# Characterizing leashes

## Lemma

*There exists an optimum homotopic Fréchet map such that the leash at every time is a shortest path (in its homotopy class).*



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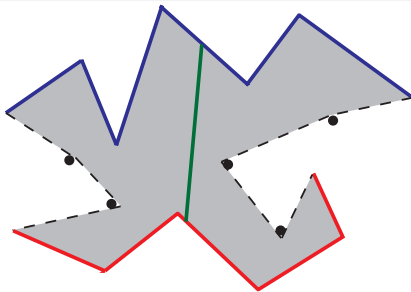
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This gives us unique shortest paths in every homotopy class, although we still have an infinite number of homotopy classes.

# Key lemma

## Lemma

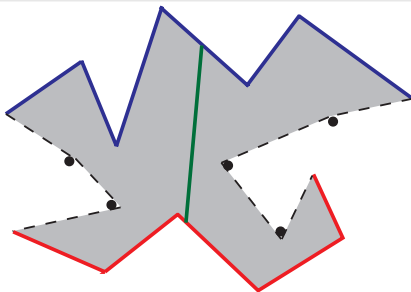
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## Lemma

*When obstacles are points, an optimal homotopy class contains a straight line segment.*

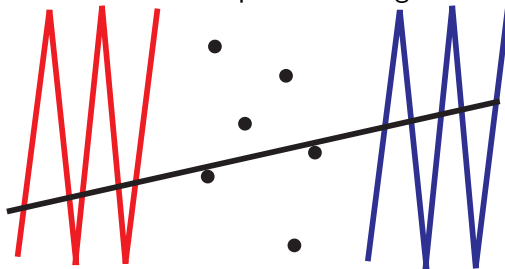


This allows us to brute force a set of possible homotopy classes which could be optimal, by trying all straight line segments.



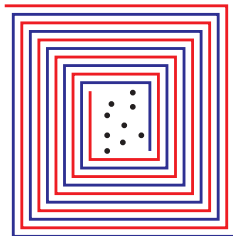
# How bad could this be?

However, there are still a lot of possible straight line segments:

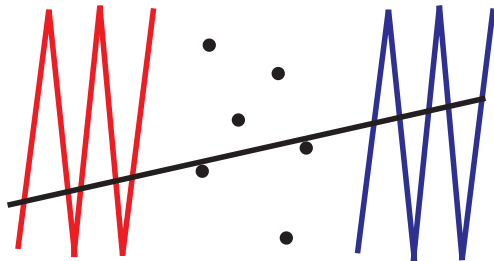


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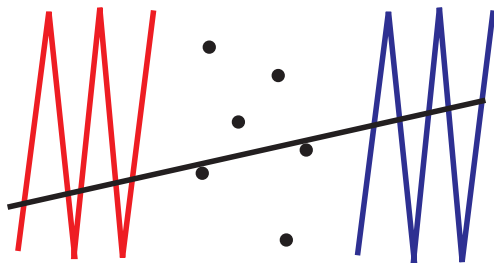


# Algorithm



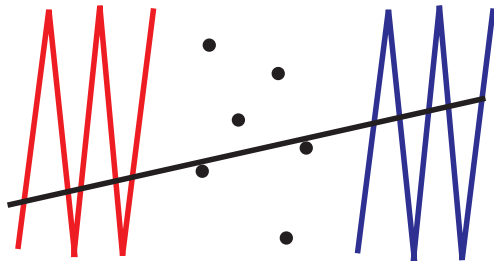
- 1 List all  $O(mn|P|^2)$  straight line segments.

# Algorithm



- 1 List all  $O(mn|P|^2)$  straight line segments.
- 2 For each homotopy class  $h$  (given by a straight line segment), compute  $F_h(A, B)$  in  $O(mn|P| \log mn|P|)$  time.

# Algorithm

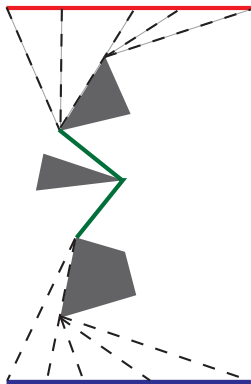


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The computation of  $F_h(A, B)$  uses techniques from the original Fréchet distance computations [Alt-Godau], as well as parametric search.

# Generalization to polygonal obstacles

Here, the optimal leash map may be **pinned** at a common subpath, which is a globally shortest path between obstacles.



This gives  $O(mn|P|^4)$  possible homotopy classes.

In addition, we must run the free space and parametric search algorithm for each relative homotopy class.

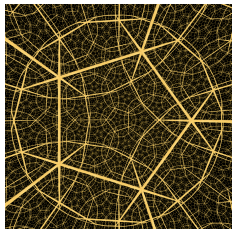
# Homotopic Fréchet Distance in other settings

It is unlikely that this approach can generalize to surfaces, since it heavily relies on nonpositive curvature. (We'll talk about weighted obstacles a bit later, though.)

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It is unlikely that this approach can generalize to surfaces, since it heavily relies on nonpositive curvature. (We'll talk about weighted obstacles a bit later, though.)

We can generalize the key lemmas to any surface of nonpositive curvature. However, the algorithmic tools in those settings are (mostly) lacking.

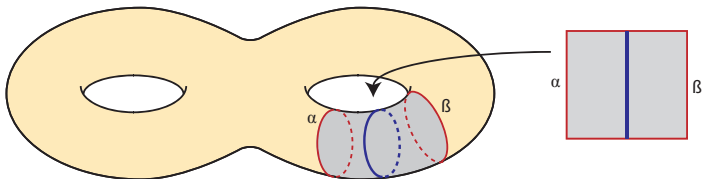




# Back to homotopy height

Recall that the height of a homotopy is an orthogonal definition to homotopic Fréchet distance:

$$d_{HH}(\gamma_1, \gamma_2) = \inf_{\text{homotopies } H} \{\sup\{|H(s, \cdot)| \mid s \in [0, 1]\}\}$$

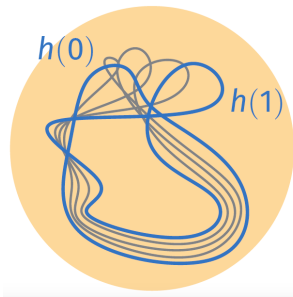


# Formalizing the notation

A homotopy through closed curves is a continuous map  $h : S^1 \times [0, 1] \rightarrow \Sigma$ , where  $\Sigma$  is a triangulated surface.

We let  $h(t)$  be the curve  $h(\cdot, t)$ , and the homotopy goes from  $h(0)$  to  $h(1)$ ; the height is then  $\sup_t \|h(t)\|$ .

An isotopy between the two curves is a homotopy where all  $h(t)$  are simple curves.



# Structure of optimal homotopies

[G. Chambers and Liokumovich] prove that some optimal homotopy is actually an isotopy.

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Sketch:

- Take a homotopy of height  $L$  from  $\gamma$  to  $\gamma'$
- Decompose into a sequence of curves  $\gamma = \gamma_1, \dots, \gamma_n = \gamma'$ , with at most 1 Reidemeister move between each  $\gamma_i$  and  $\gamma_{i+1}$



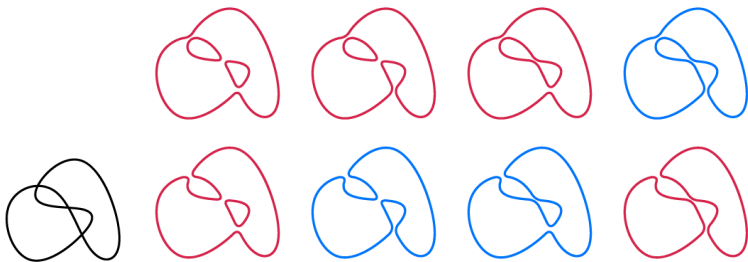
**Figure 1.1.** Homotopy moves 1 $\rightarrow$ 0, 2 $\rightarrow$ 0, and 3 $\rightarrow$ 3.

# Structure of optimal homotopies (cont)

[G. Chambers and Liokumovich] prove that some optimal homotopy is actually an isotopy.

Sketch (cont):

- We then consider all resolutions of the crossings that would get a single, simple curve

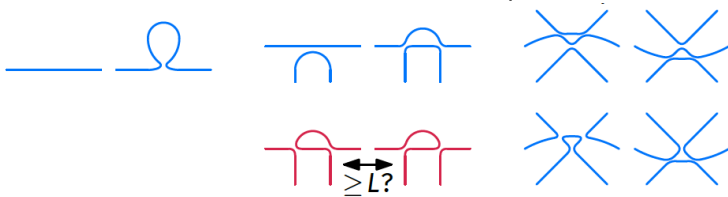


# Structure of optimal homotopies (cont)

[G. Chambers and Liokumovich] prove that some optimal homotopy is actually an isotopy.

Sketch (cont):

- Then construct “trivial” isotopies of height at most  $L$  between resolutions that are 1 Reidemeister move apart.



- Note: not all of these have a trivial isotopy between them!

# Structure of optimal homotopies (cont)

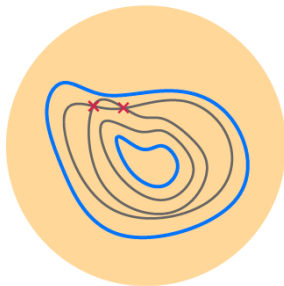
[G. Chambers and Liokumovich] prove that some optimal homotopy is actually an isotopy.

Sketch (cont):

- To fix this, they actually build a graph: vertices are the resolutions, and edges are the trivial isotopies of height  $< L$ .
- Most of the work is then proving that the graph contains a path from  $\gamma$  to  $\gamma'$ . (Surprisingly, this all boils down to the handshaking lemma.)

# Monotone isotopies

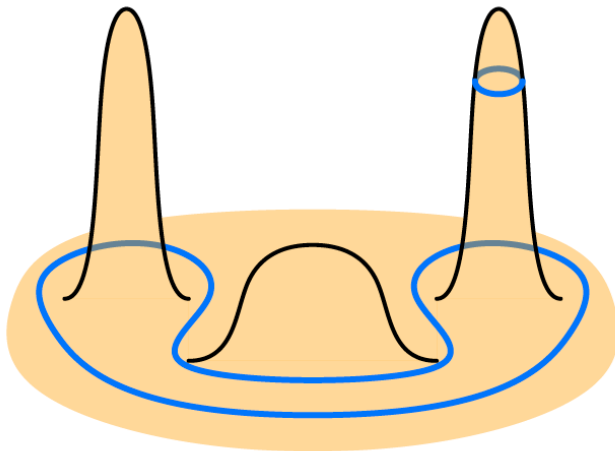
In [Chambers-Chambers-de Mesmay-Ophelders-Rotman], we show that in fact this isotopy is always monotone, so that  $h_t$  and  $h_{t'}$  are disjoint for any  $t < t'$ :





# Monotone isotopies

Note that these don't always exist! In particular, if you do not start with the boundary of the disk, the best isotopy sometimes won't be monotone:

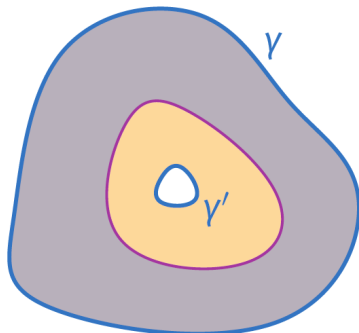
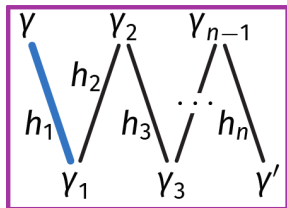


# Monotone isotopies

In [Chambers, Chambers, de Mesmay, Ophelders and Rotman] we show that monotone isotopies always exist when the curves bound an annulus.

Proof sketch:

- Decompose the isotopy into monotone sub-isotopies, where  $h_i$  goes from  $\gamma_i$  to  $\gamma_{i+1}$ :

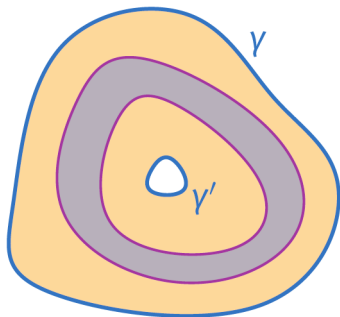
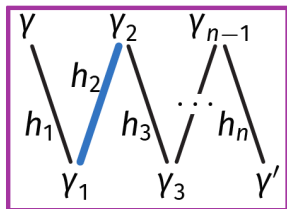


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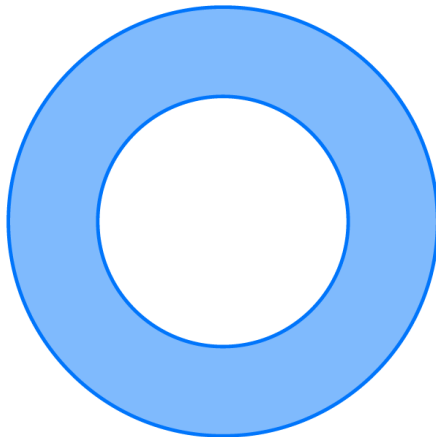
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# Monotonicity

High level idea:

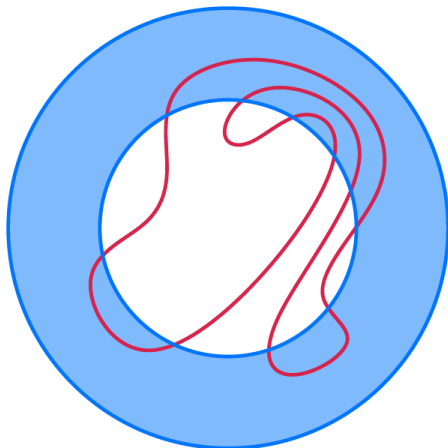
- If later parts (say  $h_{i+1}$ ) of the homotopy come back inside a previously swept portions, we want to construct a retract which stays outside:



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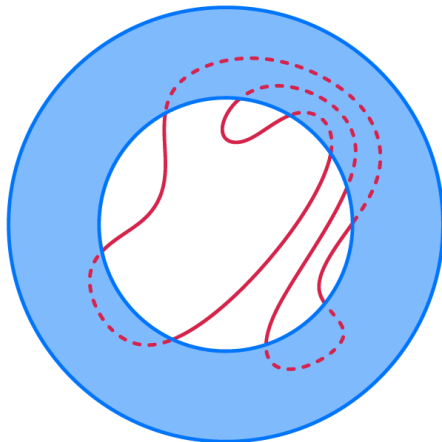
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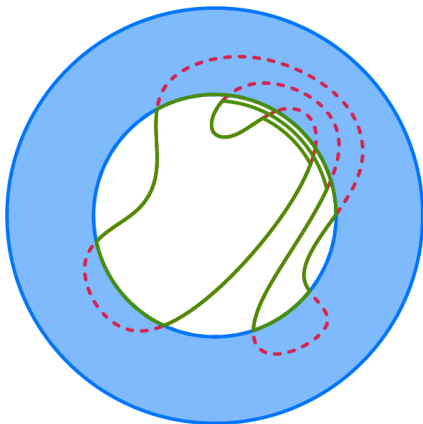
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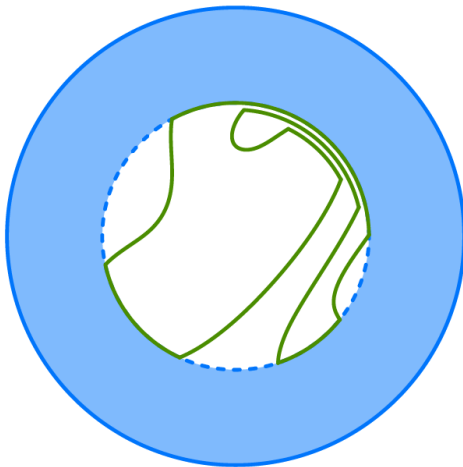
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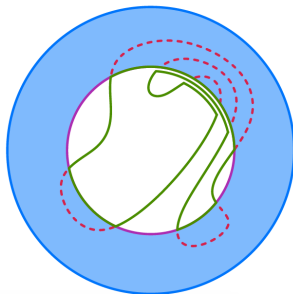




# Monotonicity

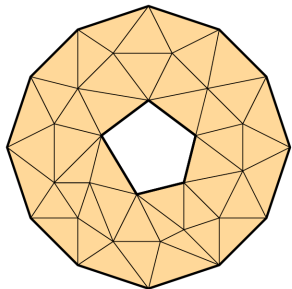
High level idea:

- These are almost isotopies of length  $< L$ : winding around can cause trouble if “bubbles” are introduced in the wrong way, and then length increases too much
- However, if the inner boundary is the shortest nontrivial curve in the annulus, we can prove that these are isotopies of length  $< L$ .

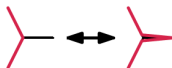


# Homotopy height is in NP

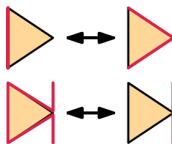
In the discrete setting, we have a triangulated annulus, and we discretize the homotopy accordingly:



Edge spike



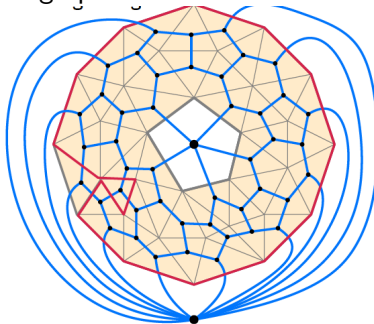
Face flip



(This is essentially what Brightwell and Winkler called a b-northward migration in their work, although they did not consider spikes.)

# Primal versus dual

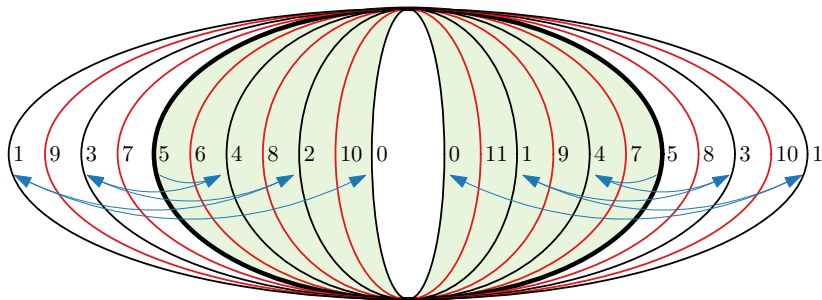
If we dualize the graph, then face moves correspond to change of crossings in the dual graph:



Monotonicity does still hold in this discretized setting if we start on the boundary of the disk, essentially since this is a very simple type of Riemannian disk.

# Non-boundary case

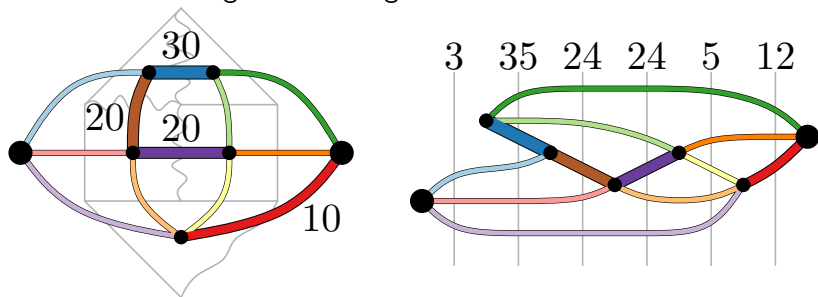
Note that we still cannot assume that the sweep is monotone if we do not begin at the boundary:



(Example courtesy of Arnaud de Mesmay)

# The dual problem

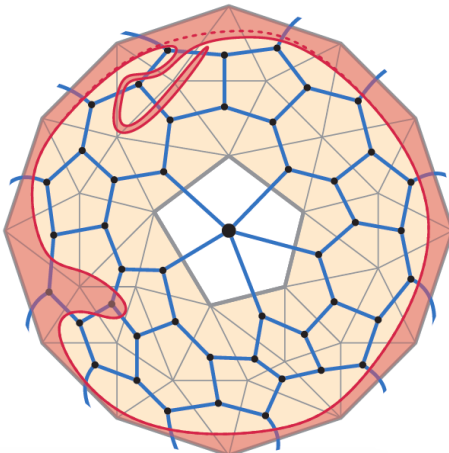
This problem in the dual is very close to the cut width of a graph, where we fix a single embedding:



Note: this is open even with unit weights, since NP hardness reductions for cut width alter the embedding of the underlying graph.

# Showing NP-Hardness

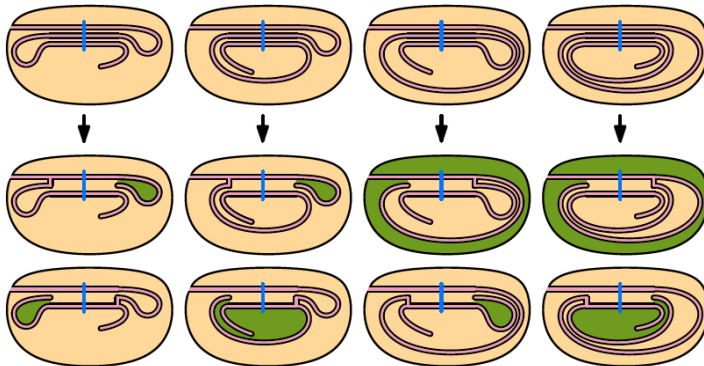
Monotonicity implies that each face flips at most once, but it does not prove the problem is in NP!



The issue is edges: those can be spiked many times from different directions.

# Bounding spikes

We show that spikes can be delayed or done early, to simplify the structure. Long paths of spikes will contain spirals, which we can simplify (essentially by case analysis):



In the end, get a quadratic bound on the number of spikes on any given edge, so homotopy height is in NP.

# Back to homotopic Fréchet

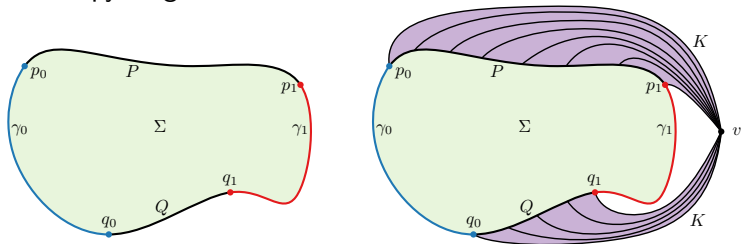
We were also able to show that a close variant of homotopic Fréchet distance is in NP as well.



# Back to homotopic Fréchet

We were also able to show that a close variant of homotopic Fréchet distance is in NP as well.

If you fix the start and end leashes of the homotopy, then you can transform an instance of the homotopic Fréchet problem into one of homotopy height:



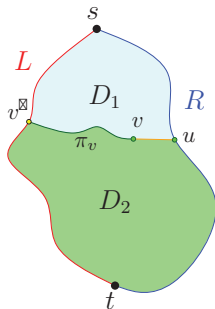
# Approximation algorithms

The first algorithmic work on homotopy height [Har-Peled-Nayyeri-Salavatipour-Sidiropoulos 2012] is  $O(\log n)$  approximation algorithm for computing both the homotopy height and the homotopic Fréchet distance between two curves on a PL surface.

# Approximation algorithms

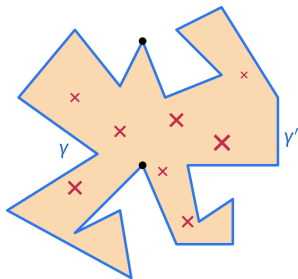
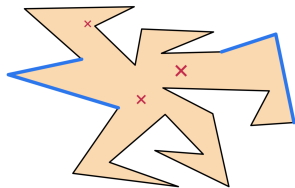
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They use a clever divide and conquer algorithm based on shortest paths for homotopy height, and then use this algorithm as a subroutine to solve homotopic Fréchet distance.



# Algorithms in some simple cases

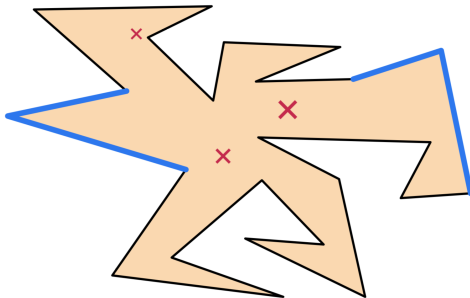
In a recent paper [Burton et al 2017], we consider homotopy height and homotopic Fréchet distance with a weighted set of point obstacles in a simple polygon, where the two input curves and initial and final leash together form the boundary of the polygon.



The leash cost at any given time in this setting is the length of the leash plus the weight of any spikes it passes over.

# Homotopic Fréchet distance in spiked polygons

It is easy to see that the optimal homotopic Fréchet map must be larger than each of the following: the initial leash, the final leash, the geodesic Fréchet distance (ignoring the obstacles), and the maximum weight point obstacle.



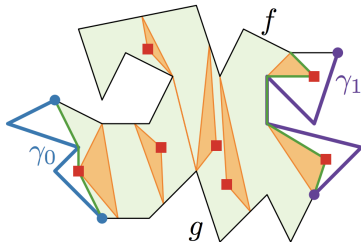
Therefore, we can compute a 2-approximation by following the geodesic Fréchet map and perturbing slightly to move over obstacles one at a time.

# Homotopy height in spiked polygons

We also solve the problem of computing homotopy height when the obstacles are unit weight points.

We prove that you can decompose the optimal homotopy in a natural way:

- fixed endpoints but can move over spikes
- endpoints of the leash can move, but don't sweep any spikes (and stay geodesic in this homotopy class)



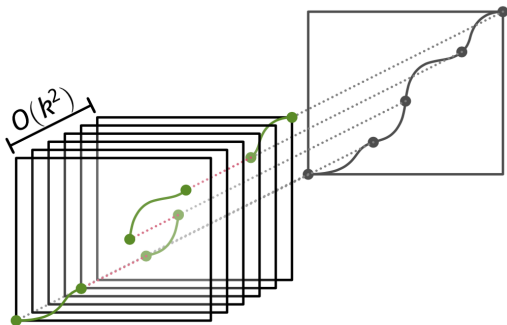
# How to compute this?

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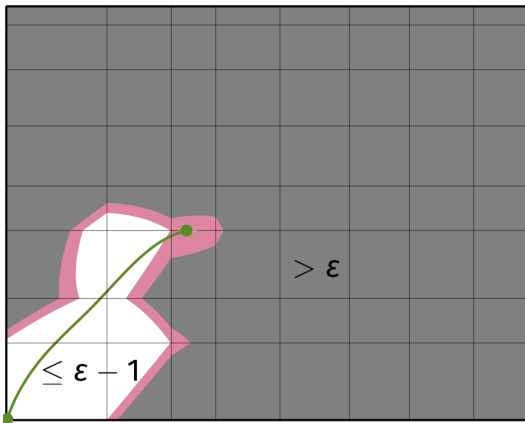
So: compute the  $\epsilon$  and  $\epsilon - 1$  free space diagrams of all relative homotopy classes which contain a straight line segment. (There are  $O(k^2)$  of these.)





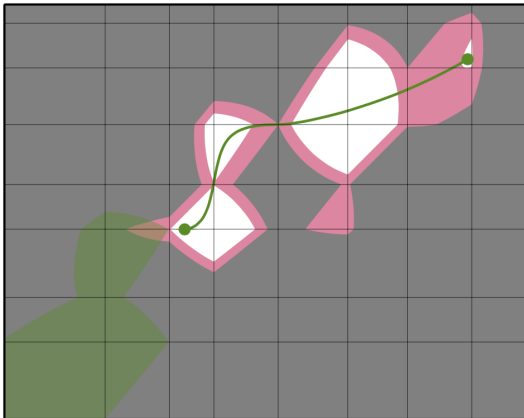
# How to compute this? (cont)

Then compute a path through these free space diagrams.



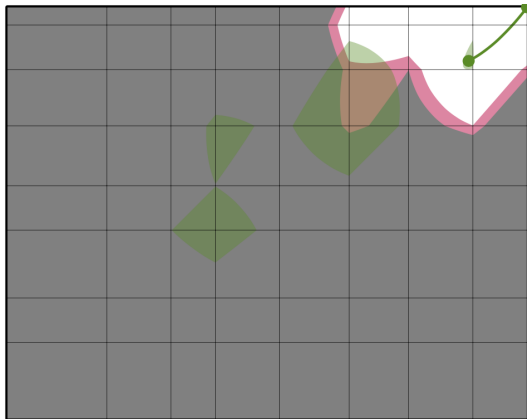
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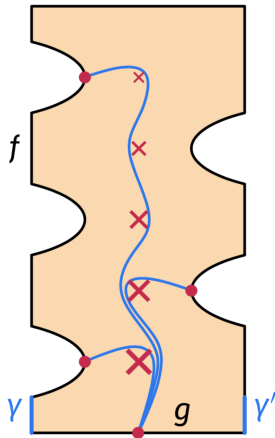


Run time:  $O(n^4 k^6 \log n + n^4 k^8 + k^{12})$ .

# Arbitrary weight obstacles

For arbitrary weight obstacles, we can't break an optimal homotopy into such nice cases, unfortunately.

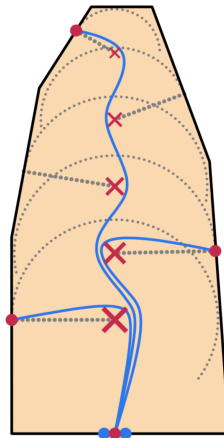
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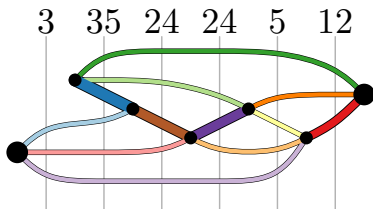
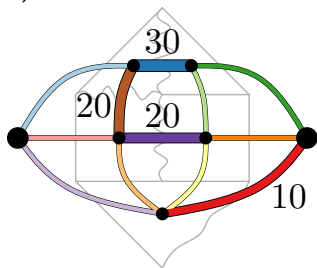
- Combinatorial versions (on graphs) are in NP.
- Some algorithms for simple obstacles in the plane. (Not very fast, though.)
- $O(\log n)$ -approximation algorithm in the plane.

However, still don't know if it's NP-Hard on graphs, or any reasonable algorithms for terrains or even arbitrary weighted point obstacles.

# Connections to other graph parameters

As mentioned earlier, homotopy height is quite naturally related to several other parameters.

Recall: homotopy height in a graph where the curve does not spike is the same as cut width of the dual graph (where embedding stays fixed):



We will call this *simple homotopy height*.



# Bar Visibility Representation

A *bar visibility representation* of a graph  $G$  is a representation where each vertex is mapped to a bar, and any two vertices are connected in  $G$  if and only if the corresponding bars have a vertical line segment that connected them and intersects no other bar.

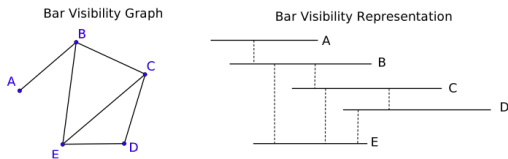


Figure 1: A bar visibility graph and its bar visibility representation.

From [Babbitt 2012]

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# Bar visibility

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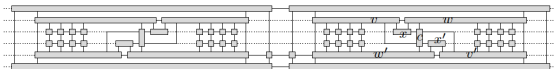
It is known that any planar graph has a bar visibility representation [Wismath 1985, Tamassia-Tollis 1986, Rosentiehl-Tarjan 1986].

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Bar visibility height is always less than or equal to 2 times the straight line drawing height: the minimum height grid such that  $G$  can be embedded on integer points and drawn with straight line edges [Biedl 2014]:



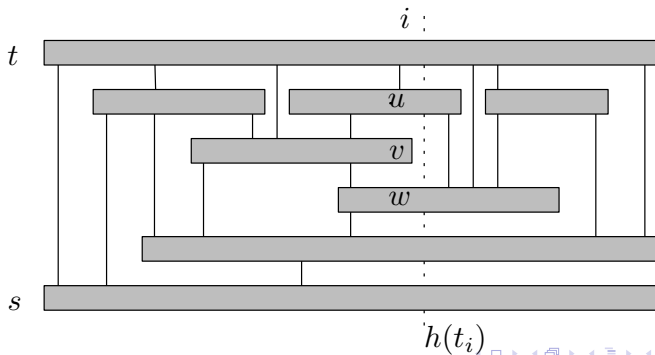
# Bar visibility and simple homotopy height

We [Biedl et. al, unpublished] also consider a new variant where we fix vertices  $s$  and  $t$  on the outer face, and ask for the minimum visibility height that places  $s$  on the top and  $t$  on the bottom of the representation.

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We prove that this is in fact exactly the same as simple homotopy height:



# Node searching or sweeping

Searching is another graph theory parameter, modeling how long it takes to sweep through a graph. In all variants, the edges of the graph are contaminated, and the graph must be cleared by guards. If at any point a cleared edge has a path to a contaminated one with no guards on the path, then it becomes recontaminated.

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- Search number: At each step, you may add a new guard to any vertex, remove a guard from a vertex, or move a guard along an incident edge of the current vertex in order to reach a new vertex. An edge is cleared when a guard moves over it.



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- Search number: At each step, you may add a new guard to any vertex, remove a guard from a vertex, or move a guard along an incident edge of the current vertex in order to reach a new vertex. An edge is cleared when a guard moves over it.
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- Connected search number: The same as node searching, but the set of edges cleared stays connected through the search.
- Monotonic search number: If the set of cleared edges only grows at every stage, then the search is monotonic.

# Connection to homotopy height

Node searching has clear connections to homotopy height: homotopies are one type of search. (This is actually why we originally looked at it, since node searching is always monotonic [LaPaugh 1993, Bienstock-Seymour 1991].)

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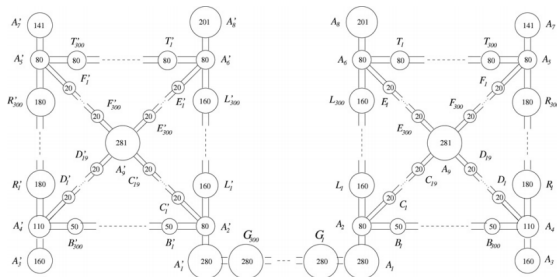
However, homotopy height is actually strictly stronger than even connected graph searching: both sides of the “cut” must stay connected for it to be a homotopy.

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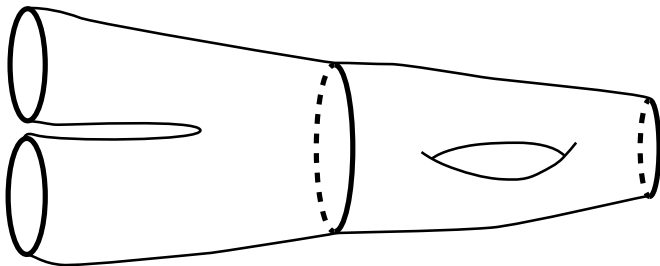
However, homotopy height is actually strictly stronger than even connected graph searching: both sides of the “cut” must stay connected for it to be a homotopy.

Interestingly, it is known that connected search number is NOT monotonic [Yang, Dyer, Alspach 2009].



# Homology “height” (or length, more accurately)

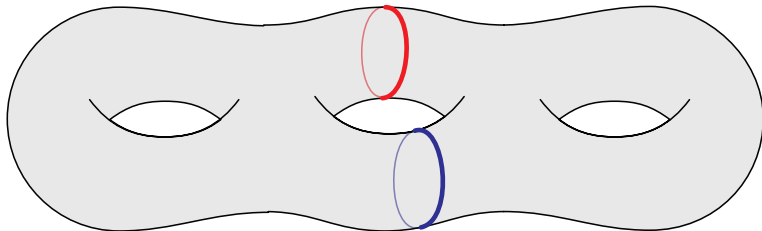
- Homology is a coarser invariant than homotopy - all homotopies produce homologies, but not all homologies come from homotopies.
- In general, homology is more tractable than homotopy - reduces to a linear algebra problem, and software is widely available and highly optimized.



# Homologous Subgraphs

## Definition

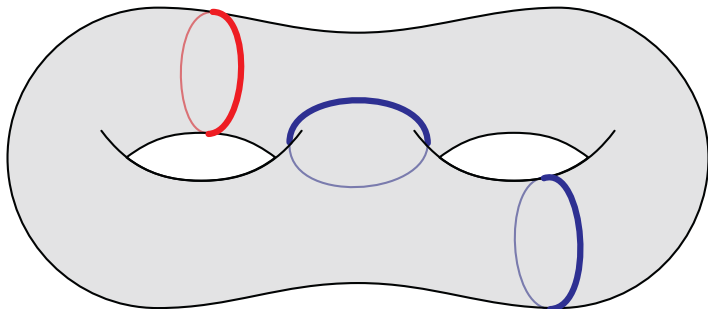
Two even subgraphs are  $\mathbb{Z}_2$ -homologous if their union forms a cut on the surface.



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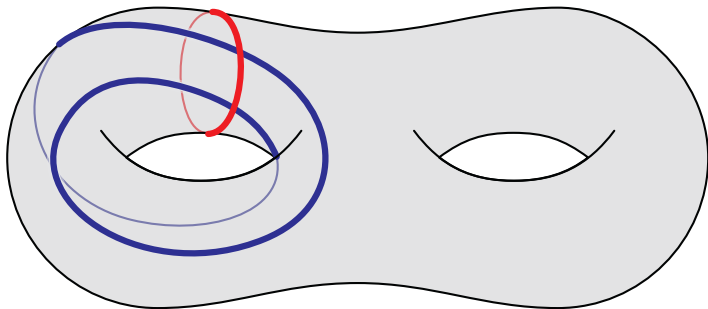




# Homologous Subgraphs

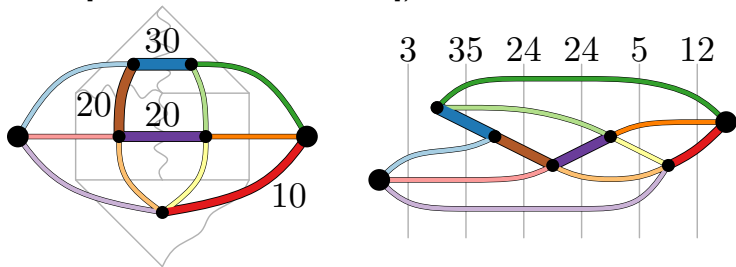
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# Homology height: NP-Hard

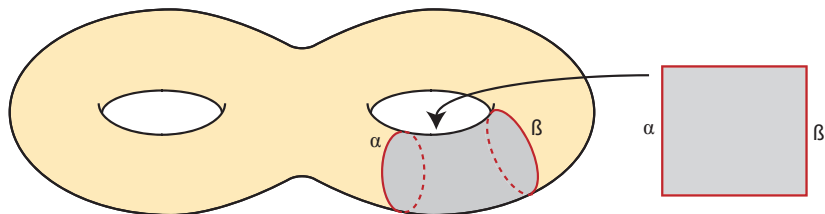
In fact, homology length is precisely the same as the cutwidth of the dual graph (once you adapt the monotonicity proof from graph searching [Bienstock-Seymour 1991]):



Here, you can line the vertices up even if they are not dual to adjacent faces: this corresponds to a new piece of the homology cycle appearing around the face, since all dual edges will be cut.

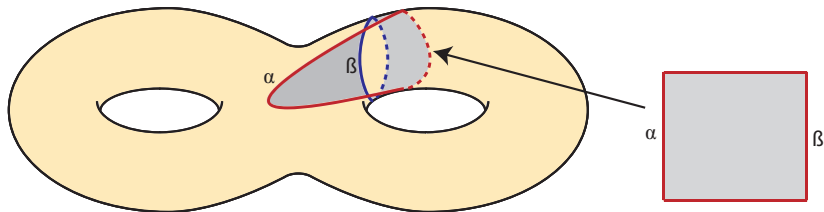
# Area of a homotopy

Recall that instead of focusing on the length or width, we can also examine the total area swept by a homotopy or homology.



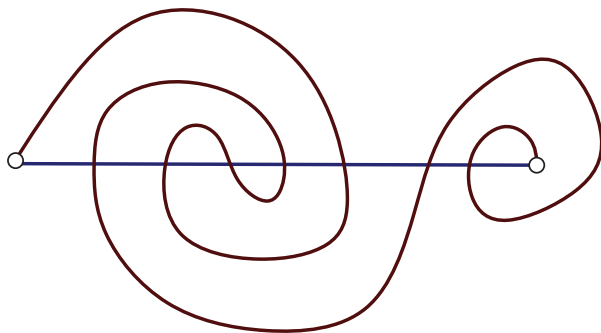
# Computing homotopy area

Surprisingly, this measure is much more tractable than any other measure based on homotopy, even for non-disjoint curves.



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More formally, given a homotopy  $H$ , the area of  $H$  is defined as:

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Note that in general, this is an improper integral, and the value for any  $H$  is not necessarily even finite.



# Douglas and Rado's work

Douglas and Rado (1930's) were the first to consider this problem, as a variant of Plateau's problem (1847) dealing with soap bubbles and minimal surfaces.

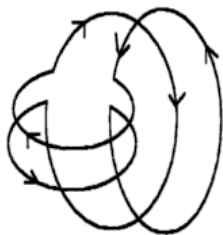


Fig. 4.1

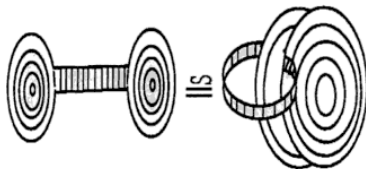


Fig. 4.2

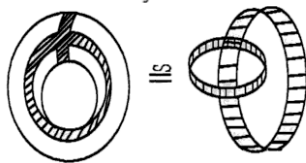
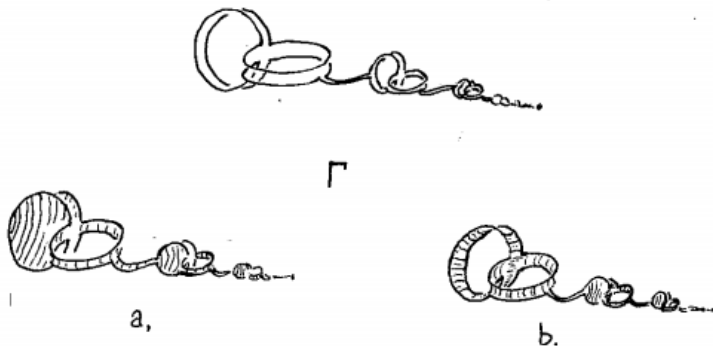


Fig. 4.3

[Minimal sub manifolds and related topics, Y. L. Xin]

# Realizing the minimum area

There is an additional problem in that to find the infimum, we might have a pathological case where a sequence of good  $H$ 's converge to something that is not even continuous.



[Lectures on Minimal Submanifolds, H. B. Lawson]

# Douglas' theorem

They developed a restricted version using Dirichlet integrals (or energy integrals) which allow control over the parameterizations of the minimal surfaces. These integrals not only minimize area, but also ensure (almost) conformal parameterizations in the space.

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## Theorem

Let  $\gamma$  be a finite Jordan curve in  $\mathbb{R}^n$ . Then there exists a continuous map  $\Gamma : \{(x, y) \in \mathbb{R}^2 : x^2 + y^2 \leq 1\} \rightarrow \mathbb{R}^n$  such that:

- 1  $\Gamma$  maps the boundary of the disk monotonically onto  $\gamma$ .
- 2  $\Gamma$  is harmonic and almost conformal
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(Well, I'm hiding a few details about the Dirichlet integrals here...)

# Necessary assumptions

In [C-Wang 2013], we consider a much simpler setting - we are either in  $\mathbb{R}^2$  or a piecewise linear surface. However, we do need some assumptions in order for the minimum area homotopy to exist.

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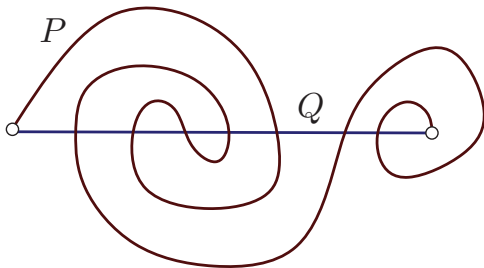
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- We must also assume the homotopy is monotone along the boundary of the domain and is regular on the interior (meaning intermediate curves are “kink-free”).
- Finally, we will assume our input curves (on  $M$ ) are simple and have a finite number of piecewise analytic components. (In practice, they will simply be PL curves.)



# Algorithm in the plane

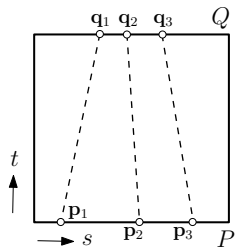
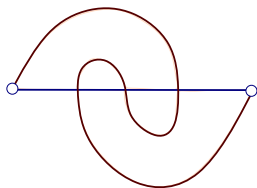
In the plane, we consider the decomposition of the plane given by the union of the two curves.



(I'm drawing continuous curves here for simplicity, but think of these as PL when we get to the running time.)

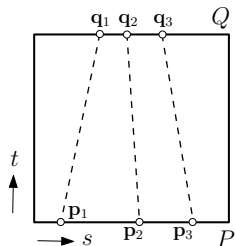
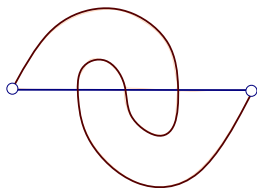
# Anchor points

Note that any vertex of intersection could either be fixed throughout the homotopy (we call this an *anchor point*) or could be moved by the homotopy.



# Anchor points

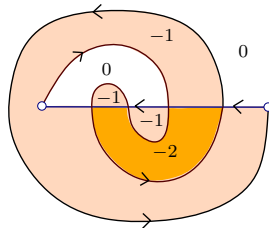
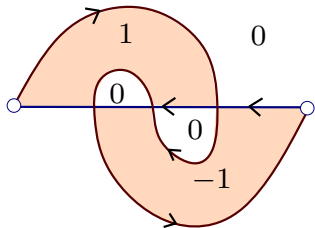
Note that any vertex of intersection could either be fixed throughout the homotopy (we call this an *anchor point*) or could be moved by the homotopy.



We prove that the ordering of the anchor points along the two curves  $P$  and  $Q$  will be the identical, and in between anchor points, we prove that the homotopy will always move locally forward.

# Winding numbers

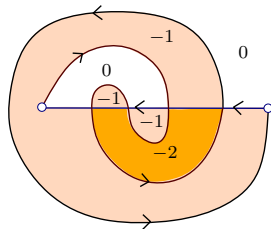
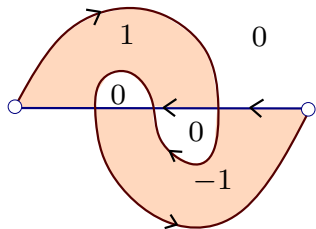
The *winding number* of a closed curve  $\gamma$  with respect to a point  $x$ ,  $wn(x; \gamma)$  is the number of times that curve travels counterclockwise around the point.



# Using the winding number

## Lemma

*Any homotopy with no anchor points will have consistent winding numbers (all non-negative or all non-positive).*

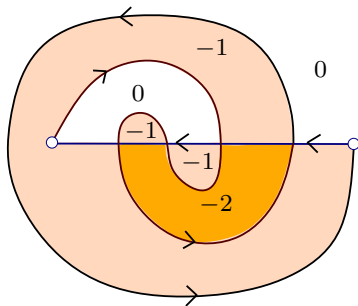


# Calculating with no anchor points

## Lemma

If  $P \circ Q$  has consistent winding numbers, then:

$$\inf_H \text{Area}(H) = \int_{\mathbb{R}^2} |\text{wn}(x; P \circ Q)| dx$$



# The algorithm: dynamic programming

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- We can compute the winding number of each planar region. If all are non-positive or non-negative, then we simply sum the areas of each region with multiplicity given by the winding number.



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- We can compute the winding number of each planar region. If all are non-positive or non-negative, then we simply sum the areas of each region with multiplicity given by the winding number.
- If the numbers are not consistent, then we know there is at least one anchor point. Since the order of the anchor points along each curve is the same, we can enumerate all the possible sets of anchor points, and in between the anchor points compute the winding numbers again.

# Running time in the plane

Let  $I$  be the number of intersections and  $n$  be the complexity of the input curves.

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# Running time in the plane

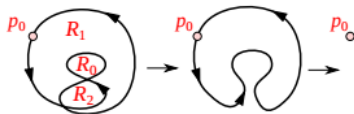
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However, this can be improved to  $O(I^2 \log I)$  time with  $O(I \log I + n)$  preprocessing if we are more careful about how we compute the winding numbers.

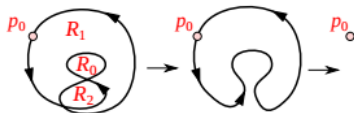
# More recent algorithms for homotopy area

There has also been recent work to compute the best area homotopy when the input curve is an immersion of a disk into the plane.



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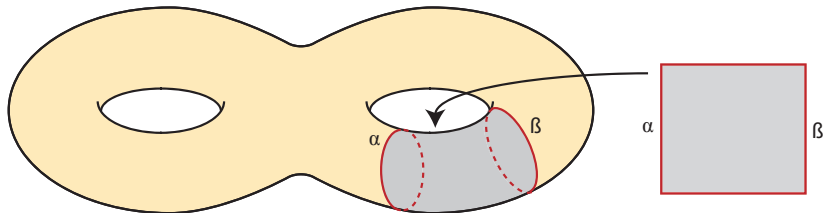
There has also been recent work to compute the best area homotopy when the input curve is an immersion of a disk into the plane.



- Nie 2014 connects this problem to the weighted cancellation norm, which is a combinatorial way to convert the best homotopy into a series of reduction moves on a word problem.
- Fasy-Karakoc-Wenk 2016 consider a different approach which is more geometric, building up an exponential time algorithm, although they are working on dynamic programming techniques to speed this up.

# Homotopy area on a surface

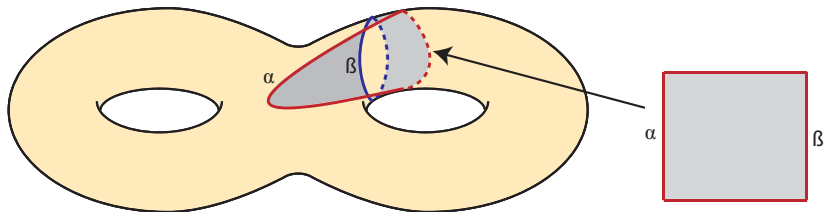
Our paper [C.-Wang] also considers the algorithm for surfaces, which builds upon the planar algorithm.



Consider two homotopic curves on a triangulated surface  $M$  with positive genus.

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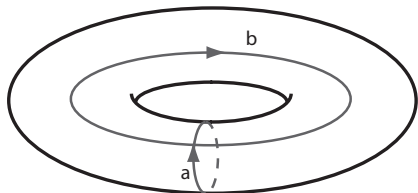


Consider two homotopic curves on a triangulated surface  $M$  with positive genus.



# The universal cover

Let  $U(M)$  be the universal covering space of  $M$ . This is a simply connected (i.e. planar) domain, along with an associated map  $\phi : U(M) \rightarrow M$  which is continuous, surjective, and a local homeomorphism.



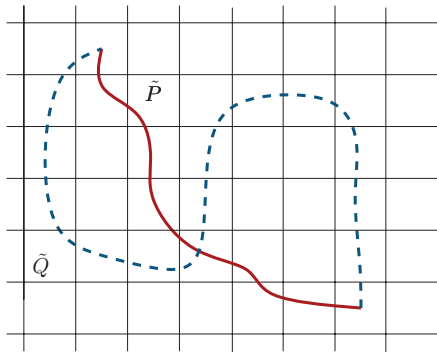
	$(-2, 2)$			$(2, 2)$	
		$(-1, 1)$	$(0, 1)$	$(1, 1)$	
		$(-1, 0)$	$(0, 0)$	$(1, 0)$	
			$(0, -1)$		
		$(-1, -1)$	$(0, -1)$	$(1, -1)$	
	$(-2, -2)$	$(-1, -2)$	$(0, -2)$	$(1, -2)$	$(2, -2)$

Arrows labeled 'a' and 'b' indicate the mapping of paths from the torus to the grid. Path 'a' is represented by vertical arrows pointing up from  $(-1, 0)$  to  $(-1, 1)$  and from  $(0, -1)$  to  $(0, 0)$ . Path 'b' is represented by horizontal arrows pointing right from  $(-1, 0)$  to  $(0, 0)$  and from  $(0, 1)$  to  $(1, 1)$ .

# Lifting $P$ and $Q$

If we fix a lift for the endpoints of  $P$  and  $Q$  in the universal cover  $U(M)$ , then  $P \circ Q$  lifts to a unique closed curve in  $U(M)$ .

Therefore, any homotopy between  $P$  and  $Q$  on  $M$  will correspond to a homotopy between their lifts in  $U(M)$  with the same area.



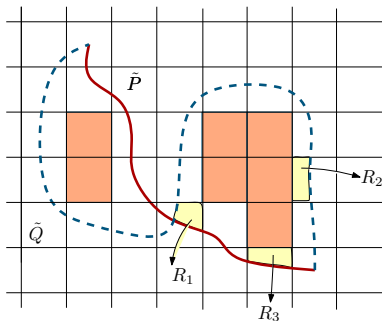
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We construct a portion of the universal cover which contains the lifts of  $P$  and  $Q$  as well as the regions inside their concatenation.

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We then use our planar algorithm in  $U(M)$ , since similar results about the winding number will hold. Since we can simplify much of the interior of the regions in our representation, the total running time here is  $O(gK \log K + l^2 \log l + ln)$ .



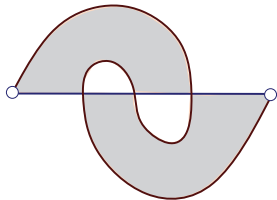
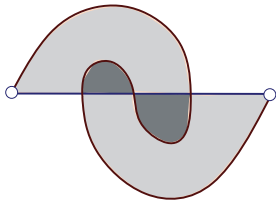
# How to compute homology area

Formally (joint work with Mikael Vejdemo Johansson, and originally considered in slightly more restricted settings in Dey, Hirani and Krishnamoorthy):

- Given cycles  $\alpha$  and  $\beta$ , try to compute  $z$  such that  $dz = \alpha - \beta$ .
- Goal: compute  $z$  with a smallest area. Recall that  $d$  is a linear operator, and  $z$  and  $\alpha - \beta$  are vectors.
- Optimization problem is then:  
 $\arg \min_z (\text{area } z)$ , subject to  $dz = \alpha - \beta$ .

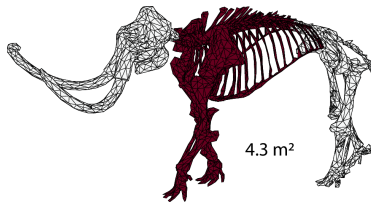
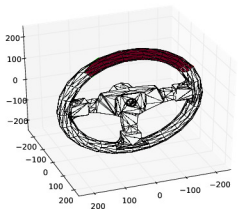
# Homology versus homotopy area

Note again that this is NOT the same as homotopy area, at least for  $d \leq 3$ :

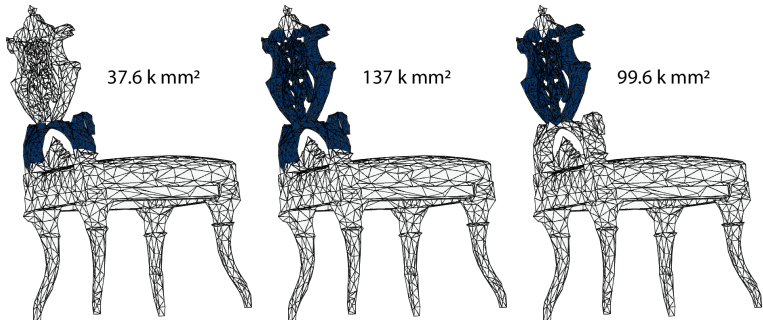


# Final algorithm for homology area

In matrix multiply time, we can compute the best area homology on meshes:

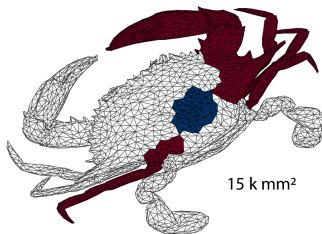
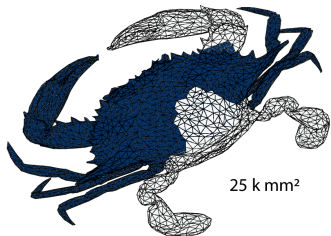


# Chair model





# Crab model



## Definition

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A *homeomorphism* is a function which is a continuous bijection where the inverse is also continuous. In our setting, this will mean that every intermediate curve in the homotopy must also have an image that is simple.

# Isotopic Fréchet Distance

In [C.-Ju-Letscher 2009], we introduced the idea of isotopic Fréchet distance:

$$\mathcal{I}(A, B) = \inf_{\substack{h : M \times I \rightarrow M \\ h(\cdot, t) \text{ homeomorphism} \\ h(x, 0) = x \ \forall x \in X \\ h(A, 1) = B}} \max_{x \in X} \text{len} h(x, \cdot)$$

In other words, what's the longest trajectory in an ambient isotopy?

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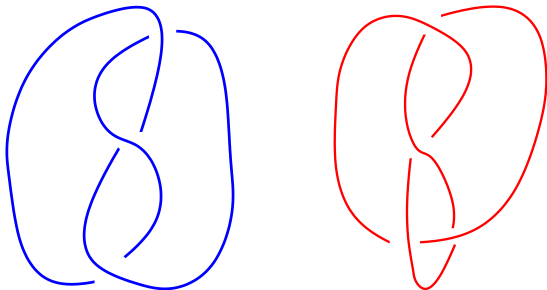
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In other words, what's the longest trajectory in an ambient isotopy?

Note the difference with homotopy height: there, the intermediate curves stayed simple, but here, we want the leashes to form an isotopy as well.

# Isotopic Fréchet Distance

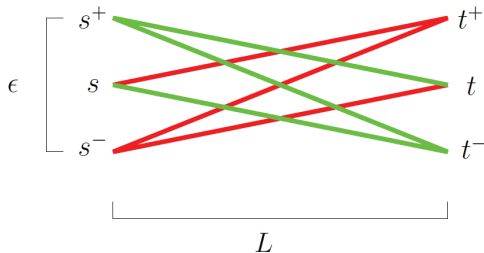
If  $A$  and  $B$  are not ambiently isotopic then  $\mathcal{I}(A, B) = \infty$ .



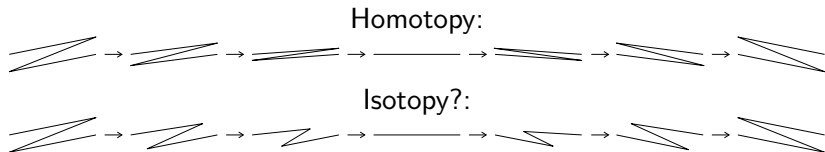
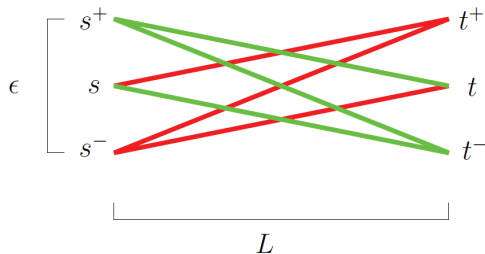
# Homotopic versus Isotopic Fréchet Distance

**Proposition** For any  $L > 0$  and  $\epsilon \in (0, L/2)$  there exists a pair of curves  $C_1, C_2 \in \mathbb{R}^2$  with

$$\begin{aligned}\mathcal{F}(C_1, C_2) = \mathcal{H}(C_1, C_2) &= \epsilon \\ \mathcal{I}(C_1, C_2) &\geq \frac{2}{9}L\end{aligned}$$



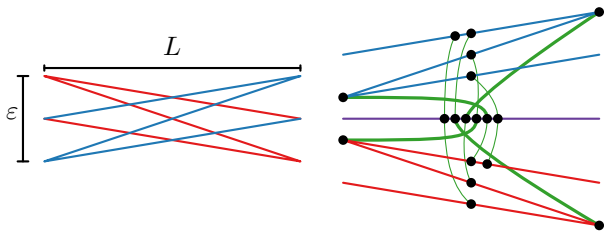
# The best homotopy versus an isotopy





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Actually, the best isotopy is even more complicated! The prior picture gave a distance of  $\sqrt{L^2 + \epsilon^2}$ . This was off by a factor of roughly 2 [Buchin-C.-Ophelders-Speckmann 2017]:





# Open questions

- There is no algorithm to compute or approximate homotopic Fréchet distance on surfaces (or even polyhedra).
- Height of a homotopy algorithms (or hardness) are also open; all that is known is an  $O(\log n)$  approximation and that it's in NP.
- These seem to be connected to all sorts of graph parameters, and even suggest some new variants to consider.
- It is unknown how to compute homotopy area between cycles on surfaces.
- Not clear if we can generalize any of these ideas (besides homology area) to surfaces instead of curves. Fréchet distance gets harder when you move to surfaces, but we don't know anything about topological variants.