5 Shortest (Homotopic) Paths

In this lecture, we'll consider a problem that combines both geometry and topology that arises in VLSI design (or at least *did* arise in VLSI design in the 1980s). Suppose we have an environment X that can be modeled as a *polygon with holes*; this is the area bounded between a simple outer polygon P_0 and several disjoint simple polygons or "holes" $P_1, P_2, \ldots, P_h$, each of which lies in the interior of P_0. (Yes, previously we used "polygon" to refer to the boundary, and now we're using "polygon with holes" to refer to the area. Welcome to mathematical jargon.)

We are also given a polygonal path π in the interior of X. Recall that two *paths* are homotopic if one can be continuously deformed to the other within X while keeping both endpoints fixed at all times. Our problem is to find the shortest path (that is, the path of minimum Euclidean length) that is homotopic in X to the given path π.

Although we could reduce to our earlier notion of homotopy by using the vertices of X as point obstacles, the pictures will be nicer (and the algorithms arguably simpler) if we use X itself as our underlying space. Our earlier formal definition of homotopy extends directly to this setting: A homotopy between two paths α and β in X is a continuous function $h: [0,1] \times [0,1] \to X$ such that the restrictions $h(\cdot, 0)$ and $h(\cdot, 1)$ are constant functions, and the restrictions $h(0, \cdot)$ and $h(1, \cdot)$ are the paths α and β, respectively.

Throughout this section, n denotes the number of vertices of the environment X, and k denotes the number of vertices in the input path π. (Yes, this is reversed from the previous lecture.) Let s ("source") and t ("target") denote the first and last vertices of π.

5.1 Shortest Paths in Simple Polygons

Let's start with the topologically trivial case where X is the interior of a simple polygon with no holes. A strengthening of the Jordan Curve Theorem due to Schönflies theorem implies that X is homeomorphic to an open circular disk. It follows that any two paths in X with the same endpoints are homotopic. Thus, we are now looking for the globally shortest path in X between the endpoints of π.

The shortest path between two points s and t in a simple polygon P is a polygonal chain, whose interior vertices lie at concave vertices of P. Imagine a taut rubber band between s to t; the rubber band will be straight everywhere, except at concave corners that it must wrap around.
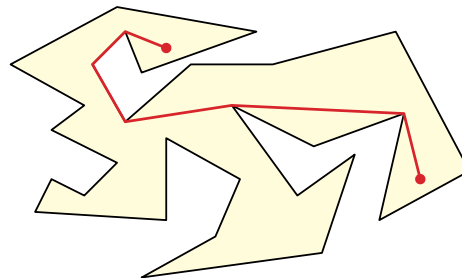

Figure 1: The shortest path between two points in a simple polygon

I'll describe an algorithm for this special case *as through* the topology of the problem were non-trivial; in particular, even though the output depends only on the endpoints of the input path π, I will still use the entire input path to guide my algorithm.

5.2 Triangulations and Dual Graphs

The first step of our shortest-path algorithm is to compute a triangulation of the polygon X. The sweepline algorithm described in the first lecture computes such a triangulation in $O(n \log n)$ time.

The *(weak) dual graph* of a polygon triangulation has a vertex for each triangle and an edge for each diagonal; two vertices are connected by an edge in the dual graph if and only if the corresponding triangles share a diagonal. We can draw the dual graph by placing each vertex at the centroid of its triangle, and drawing each edge as a polygonal path through the midpoint of the corresponding diagonal.

Lemma: *The dual graph of any triangulation of a polygon without holes is a tree.*

Proof: Suppose to the contrary that the dual graph contains a cycle C. The image of C in our drawing is a simple polygon, which I'll also call C (at the risk of confusing the reader). Any diagonal d whose dual edge is in C crosses C exactly once at the midpoint of d. It follows that one endpoint of d is inside C and the other endpoint is outside. But that's impossible, because the polygon X is connected and does not intersect the dual graph.
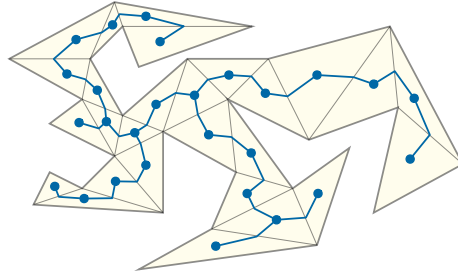

Figure 2: The dual graph of a polygon triangulation

5.3 Crossing Sequences

Next we apply the same strategy that we previously used to test whether two paths are homotopic: we compute the *crossing sequence* of π and *reduce* it as much as possible. In this setting, the crossing sequence is the sequence of diagonals crossed by π, in order along π.
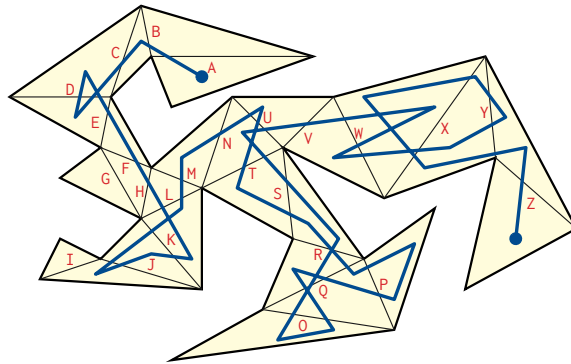

Figure 3: A path with crossing sequence ABCDDDEFHLKJJKLMNUUTSRQPPQQOOQRSTUVWWWXYYXXYZ

In our earlier homotopy-testing algorithm, we also recorded the *sign* of each crossing, but

that information is actually redundant in our current setting. Recall from our proof of the polygon triangulation theorem that any interior diagonal partitions the interior of a polygon into two disjoint subsets. Thus, if π crosses the same diagonal multiple times, those crossings must alternate between positive and negative. It also follows that we can reduce the crossing sequence by removing arbitrary adjacent pairs of *equal* symbols; for any such pair, the corresponding crossings have opposite signs. For example, the crossing sequence ABCDDDEFHLKJJKLMNUUTSRQPPQQOOQRSTUVWWWXYYXXYZ of the path in the previous figure reduces to ABCDEFHMNUVWXYZ.

We can compute the crossing sequence of π in $O(n + k + x)$ time, where $x = O(kn)$ is the length of the crossing sequence. Specifically, we find the triangle containing s by brute force; then we can repeatedly find the next crossing (if any) along the current edge of π in $O(1)$ time. Finally, we can reduce the crossing sequence in $O(x)$ time using left-greedy cancellation.

(The reduced crossing sequence of π contains precisely the edge labels that appear an odd number of times in the unreduced crossing sequence; moreover, these labels appear in the same order as their first (or last) occurrences in the unreduced crossing sequence.)

5.4 Sleeves

Let $\bar{x}$ denote the length of the reduced crossing sequence. The reduced crossing sequence defines a sequence of $\bar{x} + 1$ triangles in the triangulation of X, starting with the triangle containing the first vertex π_0 of π, and ending with the triangle containing the last vertex π_k of π. The union of these triangles is called the *sleeve* of the reduced crossing sequence.
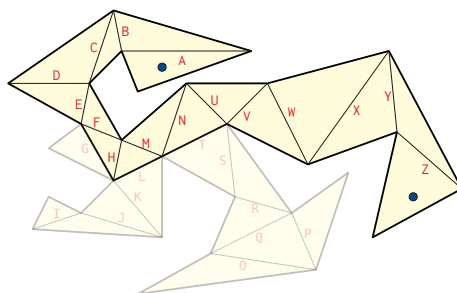

Figure 4: The sleeve of the reduced crossing sequence ABCDEFHMNUVWXYZ

Any path α in X from s to t that leaves the sleeve must cross a diagonal d on the boundary of the sleeve. The endpoints s to t lie in the same component of $X \setminus d$, so the path must cross d and even number of times. Let p and q be the first and last intersection points along α. The line segment pq is shorter than the subpath of α from p to q, so α cannot be a shortest path.

Alternatively, the crossing sequence describes a walk in the dual graph of the triangulation, starting at the vertex dual to the triangle containing s. Reducing the crossing sequence removes *spurs* from this walk—subpaths that consist of the same edge twice in a row, necessarily in opposite directions. Thus, the *reduced* crossing sequence describes a *shortest* walk—in fact, the unique simple path between its endpoints—in the dual graph. This path is also the dual graph of the induced triangulation of the sleeve.

Note that we do not actually need to *construct* the sleeve; the sequence of diagonals that the funnel crosses is exactly the reduced crossing sequence of π.

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5.5 Growing Funnels

To compute the actual shortest path from s to t, we use an algorithm independently discovered by Tompa (1981), Chazelle (1982), Lee and Preparata (1984), and Leiserson and Maley (1985). (My presentation most closely follows Lee and Preparata's.) The *funnel* of any diagonal d of the sleeve is the union of the shortest paths the from the source point s to all points on e.

The funnel consists of a polygonal path, called the *tail*, from s to a point a called the *apex*, plus a simple polygon called the *fan*. The tail may be empty, in which case s is the apex. The fan is bounded by the diagonal d and two concave chains joining the apex to the endpoints of d. The shortest path from s to either endpoint of d consists of the tail plus one of the concave chains bounding the fan. Extending the edges of the concave chains to infinite rays defines a series of *wedges*, which subdivide both the fan and the triangle just beyond d.
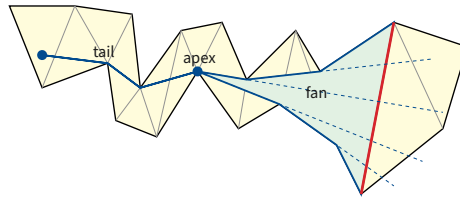

Figure 5: A typical funnel

Beginning with a single triangle joining s to the first edge in the reduced crossing sequence, we extend the funnel through the entire sleeve one diagonal at a time. Each diagonal in the sleeve shares one endpoint with the previous diagonal; suppose we are extending the funnel from diagonal pq to diagonal qr. Let o be the predecessor of p on the shortest path from s to p.

There are two cases to consider, depending on whether q and r lie on the same sides of the line through o and p or on opposite sides. We can actually detect this case in $O(1)$ time with a single orientation test.

- If q and r lie on opposite sides of line op, then the new endpoint r does not lie inside any wedge of the current fan. We can detect this case in $O(1)$ time with a single orientation test, and then extend the tunnel in $O(1)$ time by inserting r as a new fan vertex.
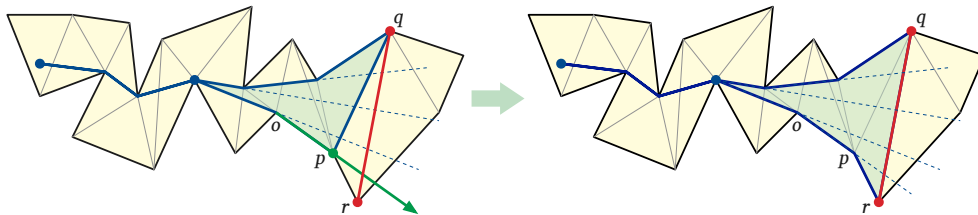

Figure 6: Growing the funnel

- If q and r lie on the same side of line op, we *contract* the funnel, intuitively by moving p continuously along the boundary edge pr. Each time the moving point crosses the boundary of a wedge, we remove a vertex from the fan. If the removed vertex is the apex, the next vertex on that side of the fan (on the shortest path from s to r) becomes the new apex. We can detect whether the moving point will cross any wedge boundary in $O(1)$ time using our standard orientation test. Thus, the total time in this case is $O(\delta + 1)$, where δ is

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the number of vertices deleted from the fan. The total number of deleted vertices cannot exceed the total number of previously inserted vertices, so the *amortized* time for this case is also $O(1)$.
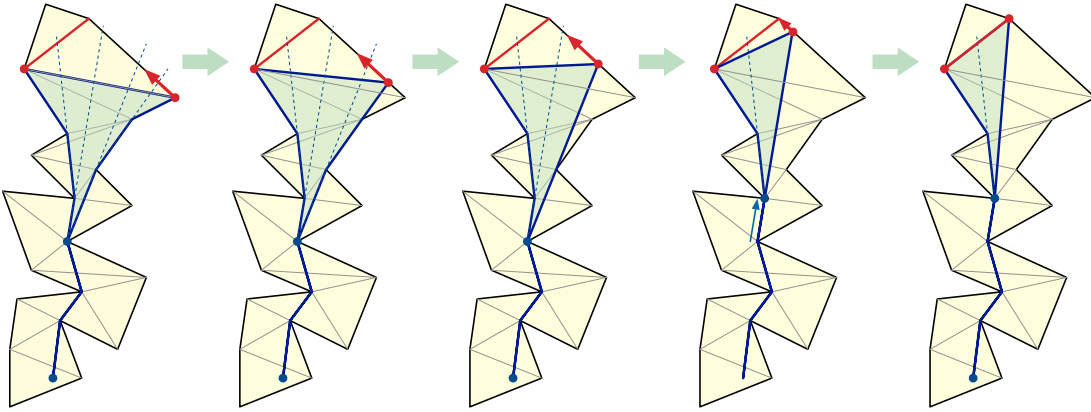

Figure 7: Shrinking the funnel

Let yz be the last diagonal in the reduced crossing sequence. To end the algorithm, we treat the line segment tz as another diagonal and extend the funnel one more time. The shortest path homotopic to π then consists of the tail of the funnel plus the concave chain of the fan containing t; we can extract this shortest path in $O(1)$ time per edge.

Summing up, we spend $O(n \log n)$ time triangulating X, then $O(k + x) = O(nk)$ time computing the crossing sequence of π, then $O(x) = O(nk)$ time reducing the crossing sequence, $O(\bar{x}) = O(nk)$ time growing the funnel, and finally $O(\bar{x}) = O(nk)$ extracting the shortest path from the final funnel.

Theorem: *Given a polygonal path π in a simple polygon X **without holes**, we can compute the shortest path in X homotopic to π in $O(n \log n + nk)$ time.*

5.6 Polygons with Holes

Now let's consider the more general case where X has one or more holes. Perhaps surprisingly, the previous algorithm needs *no modifications whatsoever* to compute the shortest path homotopic to π in $O(n \log n + nk)$ time.

- Triangulate X in $O(n \log n)$ time using the algorithm described in the first lecture. First build a trapezoidal decomposition using a sweep-line algorithm (such as Bentley-Ottmann). Then insert diagonals inside every boring trapezoid in $O(n)$ time, partitioning X into monotone mountains. Finally, triangulate these monotone mountains in $O(n)$ total time.

- Compute the crossing sequence of π with respect to this triangulation in $O(n + k + x)$ time, where x is the number of crossings. Locate the triangle containing s in $O(n)$ time by brute force, then repeatedly find the next crossing (if any) along the current edge in $O(1)$ time.

- Reduce the resulting crossing sequence in $O(x)$ time using the left-greedy reduction algorithm.

- Extend the funnel through the sleeve of the reduced crossing sequence in $O(\bar{x}) = O(nk)$ time using the standard funnel algorithm.

- Finally, extract the shortest homotopic path from the final funnel in $O(1)$ time per edge.

Theorem: *Given a polygonal path π in a simple polygon X **with holes**, we can compute the shortest path in X homotopic to π in $O(n \log n + nk)$ time.*

Same algorithm, same running time, same everything. This strikes many people as counterintuitive. After all, the sleeve can runs through the same triangle multiple times, and the funnel can self-intersect; why doesn't this cause any problems?

One way to answer this question is that *the algorithm doesn't look for self-intersections, so it's behavior can't be affected by them*. Or said differently: *the algorithm only makes local decisions, but self-intersection is a global property*. Every branch in our algorithm is based on either a comparison between two x-coordinates or an orientation test on some triple of points. As far as the algorithm is concerned, every time the funnel enters a triangle, it is entering that triangle for the very first time, or equivalently, it is entering a *new copy* of that triangle. So the sleeve of the reduced crossing sequence, while not being *geometrically* a triangulated simple polygon, is still *topologically* a triangulated simple polygon: a collection of triangles glued together along common edges into a topological disk.
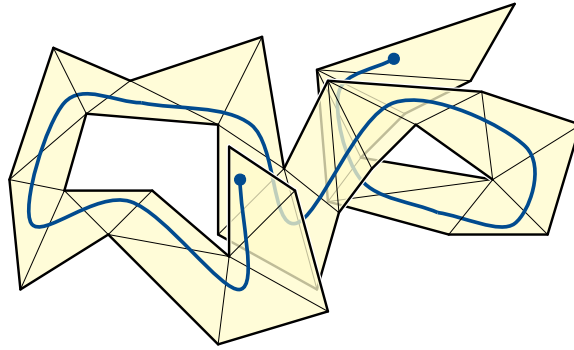

Figure 8: The sleeve of a reduced path in a polygon with holes

There is a reasonable analogy here with classical graph traversal algorithms: depth-first search, breadth-first search, and their more complex descendants. All of these algorithms maintain a set of vertices; at each iteration, each algorithm pulls one vertex v out of this set, *marks* that vertex, and then puts the *unmarked* neighbors of v into the set. Without the marking logic, unless the input graph is a tree, these algorithms will visit at least one vertex infinitely many times, each time treating it as a brand-new vertex.

In fact, even the original shortest-path algorithm does not actually require the environment X to be a simple polygon. We only require that (1) X is assembled from *any* set of Euclidean triangles by identifying disjoint pairs of equal-length edges, (2) all triangle vertices are on the boundary of X, and (3) the dual graph of the triangulation is a tree (equivalently, X is homeomorphic to a disk in the plane). More generally, the shortest-homotopic-path algorithm applies to any triangulated space satisfying conditions (1) and (2); these spaces can reasonably be called *boundary-triangulated flat surfaces*. For example, without modification, our algorithm can compute shortest homotopic paths on a triangulated Möbius band.
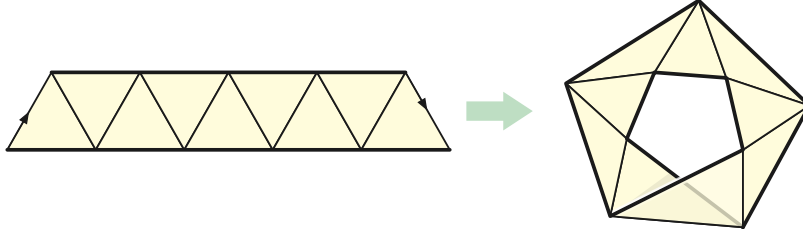
Figure 9: A flat Möbius band assembled from nine Euclidean triangles

5.7 The Universal Cover

Another explanation that may be more familiar to topologists is that our algorithm is effectively exploring the *universal cover* of the input polygon. Informally, the universal covering space of X is the infinite topological space constructed by a breadth-first search of *X without memory*.

We can define the universal cover more constructively in terms of a triangulation of X as follows. Fix a starting point s. For every reduced crossing sequence w of a path starting at s, let Δ_w denote *an independent copy* of the triangle containing the final point in that path. For example, Δ_ε is a copy of the triangle containing s. Two triangles Δ_w and Δ_x are *neighbors* if $x = wd$ for some diagonal d; glue every such pair together along their copies of d. For example, if $w = $ ABCB and $x = $ ABCBD, then we would identify the copies of edge D in Δ_w and Δ_x. We call each triangle Δ_w a *lift* of the corresponding triangle in the triangulation of X, and we call the original triangle a *projection* of Δ_w.

Figure!

Similarly, the universal cover of the dual graph, which is also the dual graph of the universal cover, has a vertex for every possible reduced crossing sequence (of a path starting at s), and an edge between two reduced crossing sequences if they differ by exactly one crossing. Unless the input polygon X has no holes, the universal cover of the dual graph is an infinite tree. The crossing sequence describes a walk from some vertex $\hat{s}$ to another vertex $\hat{t}$ in this infinite tree; removing all spurs computes the *unique* path in that tree between $\hat{s}$ and $\hat{t}$.

Figure!

5.8 Formal Definitions

Here are formal definitions for reference. A *covering map* is a continuous surjective function $p : \hat{X} \to X$, such that every point $x \in X$ has an open neighborhood U whose preimage $p^{-1}(U)$ is the disjoint union of open sets $\bigsqcup_{i \in I} U_i$, such that the restriction of the function p to each open set U_i is a homeomorphism to U. The open sets U_i are sometimes called *sheets* over U. If there is a covering space from one space $\hat{X}$ to another space X, we call $\hat{X}$ a *covering space* of X. By convention, we require covering spaces to be connected.

A space is *simply connected* if every closed curve in that space is contractible. Up to homeomorphism, every (connected) space has a unique simply-connected covering space, which is called its *universal covering space*.

The universal covering space can also be defined as the set of all *homotopy classes* of paths from

some fixed basepoint $s \in X$:

$$\tilde{X} := \left\{ [\pi] \mid \pi \colon [0,1] \to X \text{ and } \pi(0) = s \right\}$$

The covering map $p \colon \tilde{X} \to X$ maps each homotopy class $[\pi]$ of paths to their common final endpoint $\pi(1)$. Similarly, in our construction by gluing triangles, the covering map sends each triangle Δ_w to the corresponding triangle in the triangulation of X.

Concrete examples?

5.9   …and the Aptly Named Yadda Yadda

- Technicalities for point obstacles
- Bundling homotopic subpaths
- Minimum-link (homotopic) paths
- Thick non-crossing paths
- Shortest non-crossing walks / wire routing

5.10   References

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