Morphing polygonal lines: a step towards continuous generalization
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Abstract

We study the problem of morphing between two polylines that represent a geographical feature generalized at two different scales. Some cartographical generalizations are not handled well by traditional morphing algorithms, e.g., when three consecutive bends in a river or road are generalized to two bends at a smaller scale. We attempt to handle such cases by modeling the problem as an optimal matching between characteristic parts of each polyline. A dynamic programming algorithm is presented that solves the matching problem in $O(nm)$ time, where $n$ and $m$ are the respective number of characteristic parts of the two polylines. We also show the results of applying this algorithm on real road data.

1 Introduction

Visualization of geographic information in the form of maps has been established for centuries. Depending on the scale of the map the level of detail of displayed objects must be adapted in a generalization process. Be it done manually or (semi-)automatically, generalization methods usually produce a map at a single target scale. This is a well-studied field, surveyed, for example, by Weibel et al. [12].

In current geographic information systems users can interactively zoom in and out of the map, ideally at arbitrary scales and with smooth, continuous changes. However, current approaches are often characterized by a fixed set of scales or by simply zooming graphically without modifying map objects. To overcome these deficiencies continuous generalization methods are needed.

This paper studies an algorithm for continuously generalizing linear features like rivers or roads between their representations at two scales. Instead of line-simplification methods with a single target scale, we consider morphing between a source and a target scale in a way that keeps the maps at intermediate scales meaningful. Of specific interest are morphings that can deal with a certain amount of exaggeration and schematization such as reducing the number but increasing the size of road serpentines at the smaller scale. Our method first partitions the input polyline into characteristic segments and then defines distances between these segments. Based on those distances we compute an optimum morphing of the polyline segments at the two input scales using dynamic programming. We have implemented a prototype of the algorithm and compare its output with that of a simple linear morph.

2 Related work

Cecconi and Galanda [3] study adaptive zooming for web applications with a focus on the technical implementation. While maps can be produced at arbitrary scales there is no smooth animation of the zooming. A set of continuous generalization operators is presented by van Kreveld [11], including two simple algorithms for morphing a polyline to a straight-line segment.

Existing algorithms for the geometric problem of finding an optimal intersection-free geodesic morphing between two simple, non-intersecting polylines cannot be applied here because the two input polylines intersect in general. Given the correspondence between nodes of two plane graphs, Erten et al. [5] and Surazhsky and Gotsman [10] compute trajectories for an intersection-free morphing using compatible triangulations. In computer graphics, Cohen et al. [4] match point pairs sampled uniformly along two (or more) parametric freeform curves. They compute an optimal correspondence of the points w.r.t. a similarity measure based on the tangents of the curves. The algorithm is similar to ours in that it also uses dynamic programming to optimize the matching, but it does not take into account the characteristic points of geographic polylines. Samoilov and Elber [9] extend the method of Cohen et al. by eliminating possible self-intersections during the morphing.

3 Model and algorithm

In this paper, we consider the problem of morphing between two given polylines, each generalized at a different scale. Our algorithms to solve the problem can
be extended in a straightforward manner to finding a series of morphs across many scales, by solving each pair of polylines in the problem independently. The same approach can be applied to two networks with identical topology.

The problem of morphing between two polylines is two-fold. Firstly, a correspondence must be found between points on the two lines. Secondly, trajectories that connect pairs of corresponding points must be specified. Here we focus on the correspondence problem and assume straight-line trajectories.

In addressing the correspondence problem, our goal is to match parts of each polyline that have the same semantics, e.g. represent the same series of hairpin bends in a road at two levels of detail. We wish to specify that connect pairs of corresponding points must be matched up. We do this by locating points on each polyline to another. To create a morph with these desired properties, we first detect characteristic points of a polyline (Section 3.1) and use these to find an optimum correspondence (Section 3.2).

Formally, we are given two polylines $f$ and $g$ in the plane $\mathbb{R}^2$. In the correspondence problem we need to find two continuous, monotone parameterizations $\alpha : [0,1] \to f$ and $\beta : [0,1] \to g$, such that $\alpha(0)$ and $\beta(0)$ map to the first points of $f$ and $g$ and $\alpha(1)$ and $\beta(1)$ map to the last points, respectively. These two parameterizations induce the correspondence between $f$ and $g$: for each $u \in [0,1]$ the point $\alpha(u)$ is matched with $\beta(u)$.

### 3.1 Detection of characteristic points

In order to solve the correspondence problem, we first need to divide each polyline into subpolylines to be matched up. We do this by locating points on each line that are considered to be characteristic of the line; each of these characteristic points then defines the end of one subpolyline and the start of another. Previous work on generalization notes the importance of inflection points, bend points, and start and end points in defining the character of a line [8]. To find such points, we process each of the vertices in a polyline in order, checking at each if the sign of curvature has changed (an inflection point) or if the vertex is a point of locally maximal curvature (a bend point). We also apply thresholding and Gaussian filtering techniques to minimize error on noisy or poorly sampled polylines, as detailed in Algorithm 1. Gaussian filtering is a method of smoothing curves often used to assist in analyzing noisy curves; Lowe [7] gives further details and an efficient algorithm.

Algorithm 1 requires $O(|f| + n')$ time and space, where $|f|$ is the number of vertices of the polyline $f$ and $n'$ is the number of sample points. All input parameters are user-defined. Their values influence the number of characteristic points that will be detected.

### 3.2 Finding an optimum correspondence

We detect the characteristic points of $f$ and $g$ independently of each other. Assume that there are $n + 1$ such points on $f$ and $m + 1$ points on $g$, which divide the polylines into two sequences of subpolylines $(f_1, \ldots, f_n)$ and $(g_1, \ldots, g_m)$. Next, we approach the correspondence problem. Basically, there are five possibilities to match a subpolyline $f_i$:

(a) $f_i$ is mapped to the last characteristic point $g_j^\text{last}$ of a subpolyline $g_j$ (i.e., $f_i$ disappears),

(b) a subpolyline $g_j$ is mapped to the last point $f_i^\text{last}$ of $f_i$ (i.e., $g_j$ disappears),

(c) $f_i$ is mapped to a subpolyline $g_j$,

(d) $f_i$ is mapped to a merged polyline $g_{j...(j+k)}$, and

(e) $f_i$ is part of a merged polyline $f_{\ell...\ell+k}$ that is mapped to a subpolyline $g_j$.

Clearly, the linear order of the subpolylines along $f$ and $g$ has to be respected by the assignment.

Now assume that there is a morphing cost $\delta$ associated with the morph between two polylines. We suggest a morphing distance in the next section, but Algorithm 2 is independent of the concrete distance. It is based on dynamic programming and computes a minimum-cost correspondence. Algorithm 2 recursively fills an $n \times m$ table $T$, where the entry $T[i,j]$ stores the minimal cost of morphing $f_{i...i}$ to $g_{j...j}$. Consequently, we can obtain the optimum correspondence from $T[n,m]$.

The required storage space and running time of Algorithm 2 is $O(nm)$ provided that the look-back parameter $K$ is constant. Otherwise the running time...
increases to $O(nm(n + m))$. The parameter $K$ determines the maximum number of subpolyline segments that can be merged in order to match them with another segment in cases (d) and (e).

**Algorithm 2 Optimum correspondence**

**Input:** Polylines $f = (f_1, \ldots, f_n)$, $g = (g_1, \ldots, g_m)$; distance matrix $\delta$.

**Output:** Optimum correspondence for $f$ and $g$.

1. $T[0, 0] = 0$
2. $T[0, j] = T[0, j - 1] + \delta(f_1, g_j)$, $j = 1 \ldots m$
3. $T[i, 0] = T[i - 1, 0] + \delta(f_i, g_1)$, $i = 1 \ldots n$
4. for $i = 1$ to $n$ do
5. for $j = 1$ to $m$ do
6. $T[i, j] = T[i - 1, j] + \delta(f_i, g_j)$
7. $T[i, j - 1] + \delta(f_i, g_j)$
8. $\min\left(T[i, j] + \delta(f_i, g_{j+1}), g_j\right)$, $j = 2, \ldots, K$
9. $T[i - 1, j - 1] + \delta(f_i, g_j)$, $j = 2, \ldots, K$
10. $T[i - 1, j - k] + \delta(f_i, g_{j-k+1})$, $k = 2, \ldots, K$
11. $T[i - k - 1, j - k] + \delta(f_i, g_{j-k+1})$, $k = 2, \ldots, K$
12. Store pointer to predecessor, i.e., to the table entry that yielded the minimum.
13. Generate optimum correspondence from $T[n, m]$ using backtracking along pointers.

Distance measure. Algorithm 2 relies on a distance function $\delta$ that represents the morphing cost of a pair of polylines. Distance functions for polylines can be defined in many ways, e.g., morphing width [2] and Fréchet distance [1].

We define a new distance measure that takes into account how far all points move during the morphing by integrating over the trajectory lengths. Assume that two subpolylines $f_i$ and $g_j$ with uniform parameterizations $\alpha$ and $\beta$ are given. Each point $\alpha(u)$ on $f_i$ will move to $\beta(u)$ on $g_j$ along the connecting segment of length $||\alpha(u) - \beta(u)||$. Then the morphing distance is defined as

$$\delta(f_i, g_j) = \int_0^1 ||\alpha(u) - \beta(u)|| du$$  \hspace{1cm} (1)$$

and can be computed in time linear in the complexity of $f_i$ and $g_j$.

Optionally, we can add further terms to the base distance $\delta$. Adding the length difference of $f_i$ and $g_j$, or alternatively the length of the polylines $\gamma(u) := \alpha(u) - \beta(u)$ favors pairs of polylines that are roughly the same length or orientation. We can also multiply $\delta$ by the ratio of the subpolylines’ length with the total length of the containing polylines $f$ and $g$, to account for their relative visual importance.

Finally, we wish to avoid self-intersections in the morph. We do this locally by setting the effective morphing distance to $\infty$ if matching two subpolylines causes a self-intersection in the morph between them. However, in rare cases intersections between two non-corresponding subpolylines may still occur.

4 Results

We ran our implementations on a small set of French roads from the BD Carto@ and the TOP100 series maps produced by the IGN Carto2001 project [6]. For each road, we used a polyline from BD Carto@ at scale 1:50,000, and a generalized version at scale 1:100,000 from the Carto@TOP100 maps. Figures 1(a) and 1(b) show one example of a road in the dataset, at the two respective scales. The characteristic points that Algorithm 1 detected are marked by little squares. Currently, the parameters used to obtain these results were set by trial and error; so far we have no automatic process to pick reasonable values.

A sequence of snapshots of the final morph, after applying Algorithm 2, is shown in Figure 2(b). A look-back parameter $K$ of 5 was used. For the purpose of comparison, Figure 2(a) shows a simple linear morphing between the same polylines, where both polylines were uniformly parameterized to establish the correspondence between points. On a 3.0GHz Pentium 4 with 1GB RAM, the entire processing time was under 3 seconds.

The optimum-correspondence morphing shows some clear improvements over the naïve linear morphing. The linear morphing in Figure 2(a) shows one of the large serpentine sections at the top being flipped “inside-out” during the morph. In contrast, the optimum-correspondence morphing in Figure 2(b) simply expands the bends. It is evident that the total movement overall is much higher for the linear morphing than for the optimum matching morphing.

5 Concluding remarks

The algorithms in this paper should be improved in two ways. Ensuring that self-intersections do not
occurred during a morph could potentially be accomplished by utilizing the algorithm of Surazhsky and Gotsman [10] to compute non-linear trajectories for points. Also, the detection of appropriate characteristic points with little or no user interaction requires further investigation.

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References