

# MapSets: Visualizing Embedded and Clustered Graphs

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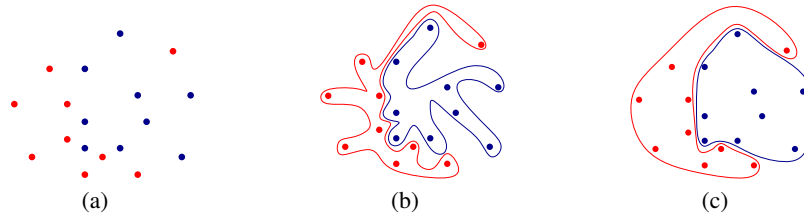
**Abstract.** We describe MapSets, a method for visualizing embedded and clustered graphs. The proposed method relies on a theoretically sound geometric algorithm which guarantees the contiguity and disjointness of the regions representing the clusters, and also optimizes the convexity of the regions. A fully functional implementation is available online and is used in a comparison with related earlier methods.

## 1 Introduction

In many real-world examples of relational datasets, groups of objects (clusters) are an inherent and important part of the input. For example, scientists belong to specific research communities, politicians are affiliated with specific parties, and living organisms are divided into biological species in the tree of life. Such clusters are often visualized with regions in the plane that enclose related objects. By explicitly defining the boundary and coloring the regions, the cluster information becomes evident. In many instances the data objects are often associated with fixed or relative positions in the plane. In geo-referenced data, for example, the positions of the objects might be based on their geographic coordinates. Thus a natural problem arises: how to best visualize graphs in which vertices are divided into clusters and embedded with fixed positions in the plane?

Several existing visualization approaches seem suitable. For example, methods for visualizing set relations over existing embedded pointsets, such as BubbleSets [7] and LineSets [2] use colored shapes to connect objects that belong to the same set. Alternatively, a geographic map metaphor can be used to represent such data. With self-organizing maps [22] or geometry-based GMaps [11], objects become cities and cluster information is captured by uniquely colored countries. While both approaches can produce compelling visualizations, we argue that neither is perfectly suited to the problem of visualizing embedded and clustered graphs.

As the number of sets increases, set-based methods generate very complex and sometimes ambiguous results. More recent methods such as KelpDiagrams [8] and KelpFusion [16] reduce visual clutter and guarantee unambiguous visualization. But more importantly, all of these methods result in overlapping regions for the sets, even when the input sets are disjoint. This unnecessarily increases visual complexity and might mislead the viewer about the disjointness of the sets. The geographic map approach suffers from a different problem. A country in the map, that represents a given cluster of vertices, might not be a contiguous region in the plane. Even though each cluster is colored with a unique color, such fragmented maps are difficult to read as human perception of color changes based on surrounding colors [19] and can be misinterpreted [13].



**Fig. 1:** (a) An embedded and clustered (red/blue) pointset. (b-c) Two different ways to construct contiguous shapes bounding points of the same color.

We want to combine the advantages of both methods, while attempting to avoid their problems. That is, we are interested in visualizing embedded and clustered graphs with non-fragmented and non-overlapping regions. While constructing such representations is easy in theory, in practice the regions may still have high visual complexity; see Fig. 1. Ideally the regions should be as *convex* as possible, as the convex hull best captures cohesive grouping according to Gestalt theory [14].

With this in mind, we describe MapSets, method for creating non-fragmented, non-overlapping regions that are as convex as possible, from a given embedded and clustered graph. We consider several criteria for measuring convexity of a given shape, and propose a novel geometric problem aiming at optimizing convexity. We include a theoretical analysis of the problem in Section 3, including a computational hardness and an approximation algorithm. Next, in Section 4, we provide a practical method for visualizing clustered graphs, which relies on the theoretical algorithm and guarantees contiguity and disjointness of the regions, and also optimizes the convexity of the regions. A fully functional implementation is available in an online system and is used in a comparison with existing techniques, which we present in Section 5.

## 2 Related Work

We review work related to the practical and theoretical aspects of the problem of visualizing embedded and clustered graphs.

**Set Visualization:** Graph clusters can be viewed as sets over graph vertices. In Venn diagrams and their generalization, Euler diagrams, closed curves correspond to (possibly overlapping) sets, and overlaps between the curves indicate intersections. Simonetto et al. [21] automatically generate Euler-like diagrams, by allowing disconnected regions, which can be complex and non-convex. Riche and Dwyer [20] propose a way to avoid the visual complexity problem by drawing simplified rectangular Euler-like diagrams, that do not depict the intersections between the sets explicitly, by duplicating objects that belong to multiple sets. In a user study, they found that it is beneficial to show intersections using simple set regions and strict containment, enabled by the duplication. For the setting where the positions of the objects is fixed, Collins et al. [7] present BubbleSets, a method based on isocontours to overlay such an arrangement with enclosing set regions. The readability of these visualizations suffer when they are many overlapping regions. LineSets [2] aim to improve the readability of complex set intersections and to minimize the overall visual clutter by reducing set regions to simple curved lines drawn through set elements. KelpDiagrams [8] incorporate classic graph-drawing “bubble and stick” style graph or tree spanners over the member points in a

set. KelpFusion [16] adds filled-in regions to provide a stronger sense of grouping for close elements. A significant limitation of all these set visualization techniques is that they produce overlapping regions even when the sets are disjoint.

**Visualizing Graphs as Maps:** The geographic map metaphor is utilized as visual interface for relational data, where objects, relations between objects, and clustering are captured by cities, roads, and countries. Using maps to visualize non-cartographic data has been considered in the context of spatialization by Fabrikant et al. [10]. Self-organizing maps, coupled with geographic information systems, render 2D maps of textual documents [23], which provide an adaptable set of tools for spatial visualization of large document collections. Maps of science showing groups of scientific disciplines are used by a wide range of professionals to grasp developments in science and technology [5]. One drawback is that self-organizing maps are very computationally expensive.

The geographic map metaphor is used in the Graph-to-Map approach (GMap) [11]. GMap combines graph layout and graph clustering, together with appropriate coloring of the clusters and creating boundaries based on clusters and connectivity in the original graph. However, since layout and clustering are two separate steps, a region representing a cluster may often be fragmented; see Fig. 7(b). Such fragmentation makes it difficult to identify the correct regions and can result in misinterpretation of the map [13].

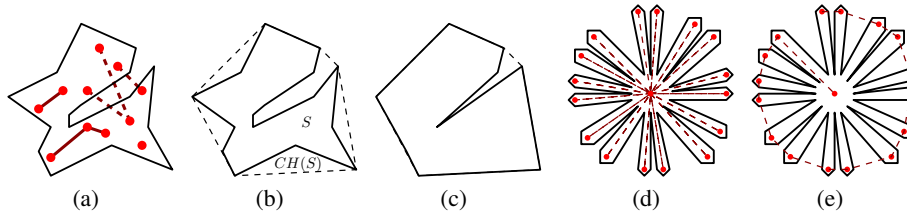
**Colored Spanning Trees:** From an algorithmic perspective, our geometric approach of optimizing convexity of regions that cover points in the plane is related to several computational geometry problems. In many problems the input is a multicolored point set, like red-blue intersection, separation, and connection problems [1,3]. Also related is the group Steiner tree problem where, for a graph with colored vertices, the objective is to find a minimum weight subtree covering all colors [17]. Also related is the problem of computing spanning graphs for multicolored point set [12]. The problem is motivated by optimizing the amount of “ink” needed to connect monochromatic points that arise when visualizing sets using the KelpFusion technique. These trees cannot be directly used as “skeletons” of regions in the plane as they can result in overlapping regions.

### 3 Constructing Contiguous Non-Overlapping Regions

We assume that the input instance consists of a set of objects  $P$  with fixed positions  $p_i \in \mathbb{R}^2$  for all  $i \in P$ , for example, cities and their geographic locations. In practical applications labels are often associated with the objects. In this case, we assume that non-overlapping bounding boxes for every label are given. The input also specifies a clustering  $C = \{C_1, \dots, C_k\}$  of the objects with  $\cup_{i=1}^k C_i = P$  and  $C_i \cap C_j = \emptyset$  for  $i \neq j$ . We wish to enclose all objects of the same cluster by a single contiguous region so that regions corresponding to different clusters do not overlap.

On one hand, simply overlaying each cluster with a convex region (e.g., bounding box or convex hull) is not always a valid solution, as it might cover elements in other clusters. On the other hand, representing clusters by some minimal regions (e.g., spanning or Steiner trees) is also not always valid, as it might result in intersecting regions.

We require regions that are contiguous and disjoint, and it is not difficult to see that such regions can be easily computed. We can begin by computing a crossing-free spanning tree of points belonging to some cluster. Once the tree is constructed, its vertices and edges become “obstacles” that should be avoided by subsequent trees. Note that all



**Fig. 2:** Convexity measures for a shape  $S$  enclosing red points. (a) Solid segments are within  $S$ , while dashed ones are not. (b) A shape  $S$  and its convex hull (dashed). (c) Area-based measure ignores boundary defects. (d-e) Ink needed to connect the points is much bigger than the length of the minimum spanning tree. The shape is enclosed in solid black, while the tree is dashed red.

the clusters will be processed as the trees do not separate the plane into more than one region. Finally, contiguous non-overlapping regions can be grown, starting from these disjoint trees. However, this procedure often generates “octopus”-like shapes that are neither aesthetically pleasant nor practically useful for visualization; see Fig. 1. Hence, we require a method for creating regions that are as convex as possible. In order to design such a method, a quality criterion for measuring the convexity of regions is needed. Next we review and formalize several convexity measures.

### 3.1 Convexity Measures

A shape  $S$  is said to be convex if it has the following property: If points  $p, q \in \mathbb{R}$  belong to  $S$  then all points from the line segment  $[pq]$  belong to  $S$  as well. The definition allows for many different ways to measure the convexity of non-convex shapes.

**Point Visibility:** For a given shape  $S$ , this convexity measure is defined as the probability that for points  $p$  and  $q$ , chosen uniformly at random from  $S$ , all points from the line segment  $[pq]$  also belong to  $S$  [25]. The result is a real number from  $[0, 1]$ , with 1 corresponding to convex shapes.

A problem with this definition is that it is difficult to compute, even if  $S$  is a polygon. Therefore, we consider its discrete variant, taking into account that the input of our problem specifies points in the plane; see Fig. 2(a).

**Vertex Visibility:** This measure takes into account how many segments  $[pq]$  are completely in  $S$  for pairs of input points  $p, q \in P$  of the cluster corresponding to  $S$ . The measure is defined as  $\frac{\sum_{p,q \in P} \delta(p,q)}{|P|^2}$ , where the sum is over all pairs of input points  $P$  and  $\delta(p, q) = 1$  if  $[pq]$  lies inside  $S$  and  $\delta(p, q) = 0$ , otherwise. The result is a real number from  $[0, 1]$ , with 1 corresponding to convex shapes.

**Convex Hull Area/Perimeter:** Recall that the smallest convex set which includes a shape  $S$  is called the *convex hull*  $CH(S)$  of  $S$ ; see Fig. 2(b). The area-based convexity measure is defined as  $\frac{Area(S)}{Area(CH(S))}$ ; it is frequently used and appears in textbooks [24]. The result is a real number from  $[0, 1]$ , with 1 corresponding to convex shapes. Unlike visibility-based measures, the convex hull-based one is very easy to calculate efficiently and is robust with respect to noise. However, the definition does not allow to detect defects on boundary that have a relatively small impact on the shape area; see Fig. 2(c). The perimeter-based definition attempts to remedy this:  $\frac{Perimeter(S)}{Perimeter(CH(S))}$ .



**Fig. 3:** (a) An input for CST with  $n = 10$  points and  $k = 3$  colors. (b) An optimal solution with minimum ink containing Steiner points.

If a shape  $S$  is convex, then there exists a minimum spanning tree on the given point set such that every edge of the tree lies completely in  $S$ . On the other hand, non-convex shapes do not necessarily admit such a spanning tree. Hence, the length of a shortest curve that belongs to  $S$  and connects all the input points is an indicator of convexity of  $S$ . In the following measure, we compare the length of such a curve (or equivalently, the amount of “ink” needed to connect all the points) with the length of a minimum spanning tree on the same point set; see Figs. 2(d)-2(e).

**Minimum Ink:** Let  $\text{INK}(P)$  be the length of the shortest curve connecting all vertices of  $V$  lying in  $S$ , and let  $\text{MST}(P)$  be the length of the minimum spanning tree of  $V$ . The measure is defined as  $\frac{\text{MST}(P)}{\text{INK}(P)}$ . Here, 1 indicates the best possible value (though, it does not always correspond to a convex shape), while smaller values are worse.

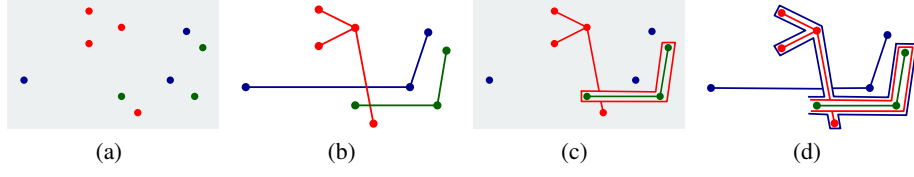
Note that there are advantages and disadvantages of all of the proposed convexity measures, and there are also many other ways to define convexity of shapes or polygons. In an attempt to balance theoretical and practical considerations, we focus on visibility-based and the ink-based measures. Note that similar ink-based criteria are used to measure the “simplicity” of shapes used for constructing LineSets and Kelp-Diagrams. By minimizing the ink needed for drawing, all of these techniques aim to reduce visual clutter and increase the readability of the representation.

### 3.2 Algorithm for Ink Minimization

Here we study a problem motivated by computing contiguous regions with minimum ink. The input consists of  $n$  points in the plane, and each point is associated with one of  $k$  colors. The CST (COLORED SPANNING TREES) problem is to connect points of the same color by mutually non-intersecting curves of shortest total length. It is easy to see that in an optimal solution each curve forms a tree spanning points of the corresponding color. In general, the trees may use additional (Steiner) points that do not belong to the original pointset; see Fig. 3.

Computing an optimal solution for CST is NP-hard. This follows directly from the observation that the known NP-complete MINIMUM STEINER TREE problem is a special case of CST, in which the input consists of monochromatic points. Next we present a heuristic for CST and prove that it is an approximation algorithm in the theoretical sense, and hence produces solutions guaranteed to be close to the optimum.

We refer to the minimum spanning tree of a set of points  $P$  as  $\text{MST}(P)$ . The minimum Steiner tree of the points is referred to as  $\text{SMT}(P)$ . Slightly abusing notation, the lengths of the trees are also denoted by  $\text{MST}(P)$  and  $\text{SMT}(P)$ . We use the Steiner ratio, denoted by  $\rho$ , which is the supremum of the ratio of the total length of a minimum spanning tree to the total length of a minimum Steiner tree. It is conjectured that



**Fig. 4:** Steps of the algorithm for the CST problem. (a) An input with  $n = 10$  points and  $k = 3$ . (b) Computing minimum spanning trees. (c) Bounding the tree having the shortest length, and removing red-green crossings. (d) Merging with the blue tree.

$\rho = \frac{2}{\sqrt{3}} \approx 1.15$ , but the conjecture is still open. Chung and Graham [6] showed a bound of  $\approx 1.21$ , which is the best-known upper bound on  $\rho$ .

We begin with description of our algorithm in the setting when the input consists of blue and red points. In the first step, we compute a minimum spanning tree of the blue points (ignoring the red ones), and a minimum spanning tree of the red points; see Fig.4(b). If the trees do not intersect, then they form a solution for CST. Otherwise, we create a red “shell” bounding the blue tree; see Fig.4(c). Note that now all red-blue crossings appear inside the constructed shell. To eliminate the crossings, we remove all portions of the red tree inside the shell; the operation clearly keeps the red tree connected. Finally, the red curve, consisting of the original spanning tree and the constructed shell, can be transformed to a tree by disconnecting its cycles; see Fig.4(d).

The general algorithm works in the following steps. First, create a minimum tree  $\text{MST}(C_i)$  spanning the set of points  $C_i$  for  $1 \leq i \leq k$ , ignoring points of the other colors. Sort the colors with respect to the length of the corresponding spanning trees. Without loss of generality, we may assume that the resulting order is  $C_1, \dots, C_k$  and  $\text{MST}(C_1) \leq \dots \leq \text{MST}(C_k)$ . Then the resulting curve for  $C_1$  is the tree  $\text{MST}(C_1)$ . A curve for each successive color  $C_i$  is constructed by adding a “shell” bounding the curve corresponding to  $C_{i-1}$ . Note that the length of the shell is exactly  $2 \sum_{j < i} \text{MST}(C_j)$ , since it bounds all the spanning trees corresponding to already processed colors; see Fig. 4. The length of a curve for  $C_i$  is then  $\text{MST}(C_i) + 2 \sum_{j < i} \text{MST}(C_j)$ .

In order to analyze the algorithm, we denote the amount of ink in the optimal solution by  $\text{OPT}$ , and the total length in the constructed solution by  $\text{ALG}$ . It is easy to see that an optimal solution induces a curve connecting all points of the same cluster, that is, the solution is a Steiner tree for the set of points (but not necessarily the minimum one). Hence,  $\text{OPT} \geq \sum_i \text{SMT}(C_i)$ . On the other hand,

$$\text{ALG} \leq \sum_i (\text{MST}(C_i) + 2 \sum_{j < i} \text{MST}(C_j)) = \sum_{i=1}^k (2k - 2i + 1) \text{MST}(C_i).$$

Hence, we have

$$\begin{aligned} \frac{\text{ALG}}{\text{OPT}} &\leq \frac{\sum_{i=1}^k (2k - 2i + 1) \text{MST}(C_i)}{\sum_{i=1}^k \text{MST}(C_i) / \rho} = \\ &= \rho \frac{\sum_{i=1}^{\lfloor k/2 \rfloor} (2k - 2i + 1) \text{MST}(C_i) + \sum_{i=1}^{\lceil k/2 \rceil} (2i - 1) \text{MST}(C_{k-i+1})}{\sum_{i=1}^k \text{MST}(C_i)} \leq k\rho. \end{aligned}$$

Hence, our algorithm is a  $(k\rho)$ -approximation for the CST problem for any  $k \geq 1$ .

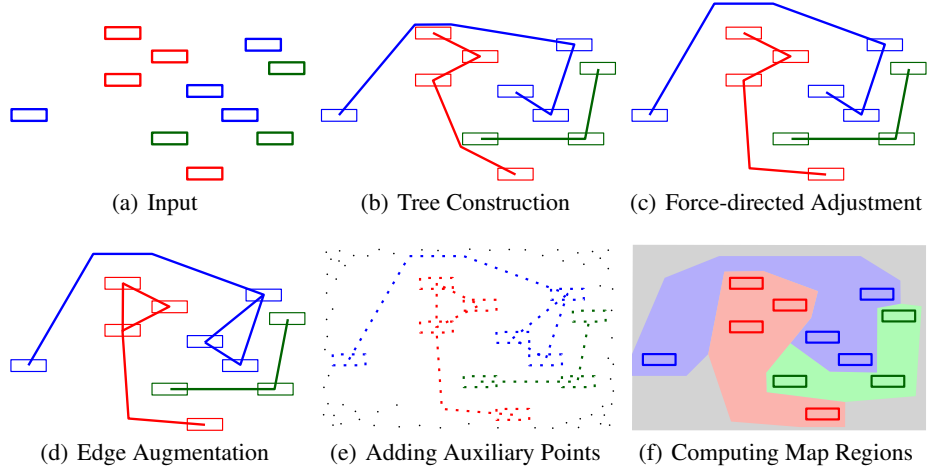


Fig. 5: Algorithmic pipeline of MapSets.

## 4 MapSets

Here we describe MapSets, starting with a high-level overview; see Fig. 5. We assume that the input is a set of rectangular shapes (bounding boxes of labels) embedded in the plane along with a clustering. In the first step, we compute spanning mutually non-crossing trees interconnecting centers of rectangles corresponding to the same cluster, while minimizing the total ink needed to draw the trees. In the second step, we modify the trees by adding buffers of free space around the segments of the trees, using a force-directed heuristic. In the third step, we try to optimize the convexity of the resulting regions based on the vertex visibility measure, by adding edges between vertices in the same cluster, while ensuring that edges of different clusters do not cross. In the fourth step, we use the modified trees and added edges to build contiguous non-overlapping boundaries for all clusters.

**Tree Construction:** In order to construct the trees, we use the approximation algorithm described in Section 3.2. For each cluster, we first compute a minimum tree spanning the set of rectangle centers, ignoring other clusters. The clusters are then sorted in non-decreasing order by the length of the computed trees and processed in this order. At each step we consider all the precomputed trees as obstacles that should be avoided when constructing the current tree. The rectangles are also treated as obstacles. We compute a sparse visibility graph on the set of obstacles, where the vertices are all the centers and corners of the rectangles, and there is an edge between two vertices if one can draw a straight-line segment without crossing the obstacles. The sparse visibility graph (unlike the full visibility graph) has a linear number of edges and can be constructed efficiently [9]. We then compute shortest paths (of the visibility graph) between every pair of rectangles of the current cluster. From these shortest paths, we compute a minimum spanning tree for the current cluster. We add the tree to the set of obstacles and proceed with the next cluster.

**Force-directed Adjustment:** This step improves the constructed trees. Our goal is to provide some free space around the edges of the trees so as to avoid (1) narrow

channels between parts of the same country and (2) country borders lying too close to the input vertex labels. To accomplish this, we consider an adjustment graph  $H$  in which vertices are the end points and bends of the constructed trees and edges are maximal straight-line segments of the trees. We then build a force system moving the vertices of  $H$  that correspond to the bends of the tree. The system relies on the following forces.

- **Vertex-vertex attraction.** We would like to keep the ink of the drawing low. Therefore, for every vertex of  $H$ , there is a force pushing the vertex towards its neighbor vertices in  $H$ .
- **Edge-edge repulsion.** This repulsive force attempts to push the edges of  $H$  apart to provide enough space to draw the regions. In order to compute the force, it is convenient to replace edges of  $H$  with cylinders of a specified thickness. Then, if two cylinders corresponding to different trees intersect, the force repels them away from each other. This force also ensures that the trees do not overlap and do not intersect during the adjustment process.
- **Edge-label repulsion.** This force prevents edges from being routed too close to the input text labels. Again, it is convenient to consider the edges of  $H$  as cylinders. If a cylinder occludes a label, then we introduce a repulsive force moving the corresponding vertices of  $H$  away from the label.

We use iterative refinement similar to that used in drawing graphs with edge bundles [4] to adjust the positions of the vertices of  $H$  under these three forces: repulsive forces have equal priorities, and the attractive force is weaker. In our experiments, the force system provides the desired buffer of free space around the trees and converges quickly; see Fig. 8.

**Edge Augmentation:** In this step we try to optimize the convexity of the regions using the vertex visibility metric. Consider all possible straight-line segments connecting centers of rectangles corresponding to the same cluster. Our goal is to select and add as many of these segments as possible, subject to the condition that they do not cross each other. To this end, we construct a graph  $H$  in which vertices are the straight-line segments. A segment is added to  $H$  only if it does not intersect the trees found in the previous step. Two vertices of  $H$  are connected by an edge if the corresponding straight-line segments cross each other. Notice that now the problem reduces to the problem of finding a maximum non-crossing (independent) set of segments in the plane. The problem can be solved optimally in polynomial time for two clusters, that is, if  $k = 2$ . Indeed, in this setting the graph  $H$  is bipartite, and the size of a maximum independent set in a bipartite graph equals to the number of edges in a minimum edge covering by König’s theorem. The latter can be found using a maximum matching algorithm. Unfortunately, the general variant is NP-hard even for  $k = 3$  [15]. Therefore, unless  $k = 2$ , we use a greedy strategy to solve the problem. At every step, we choose the minimum degree vertex in  $H$  and remove its neighbors. It is well-known that this strategy guarantees an approximation ratio of  $(\Delta + 2)/3$  on graphs with maximum degree  $\Delta$ .

**Adding Auxiliary Points and Computing Map Regions:** Given the initial placement of the labels and curves connecting the labels from the previous steps, we want to create a map: that is we need explicit regions grouping together labels and curves in the same cluster. A naive method is to form the constrained Voronoi diagram of the labels and computed curves. However, this often results in sharp corners and angular outer



boundaries of the constructed regions. Hence, we generate more natural boundaries by adding dummy points to the current embedding, as in the GMap framework [11]. There are three types of the dummy points. First, random points, sufficiently far away from the set of the input labels, lead to more rounded and thus more realistic region boundaries. Second, random points along bounding boxes of the labels help ensure that the labels are drawn inside the countries. Finally, auxiliary points are added along all the edges constructed on the previous step. These points are important in our algorithm, as they keep the regions connected. The distance between consecutive points on an edge is chosen to be less than the distance to any other point of a different color. After adding the dummy points, we compute the Voronoi diagram of the set of all points. Voronoi cells that belong to the points of the same color are merged together to produce the final map.

### Time Complexity

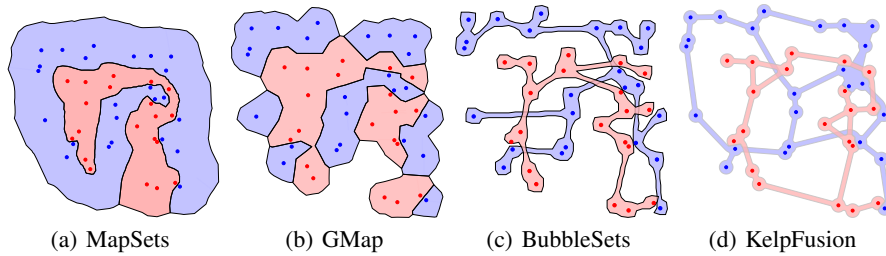
Now we discuss the complexity of our algorithm on an input with  $n$  points and  $k$  clusters, assuming we can compute distances and intersections between geometric primitives (points, line-segments, rectangles) in constant time. The sparse visibility graph can be constructed in  $O(n \log n)$  time and it contains  $O(n)$  edges [9]. Therefore, computing all pairwise distances takes  $O(n^2)$  time and finding a minimum spanning tree for one cluster takes  $O(n^2 + n \log n)$  time. Summing over all clusters, we get  $O(kn^2)$ . In the iterative force-directed heuristic we compute forces between pairs of edges, which can take  $O(n^2)$  in the worst case. Hence, the time complexity of the force-directed heuristic is  $O(cn^2)$ , where  $c$  is the maximum number of iterations in the adjustment ( $c = 10$  in our implementation). The complexity of the edge augmentation step is  $O(n^3)$ , as we may add quadratic number of edges in the greedy process. Finally, computing the boundaries takes  $O(n \log n)$  time. Therefore, the overall time complexity is  $O(kn^2 + n^3)$ . More details and actual running times are given in the next section.

## 5 Experiments

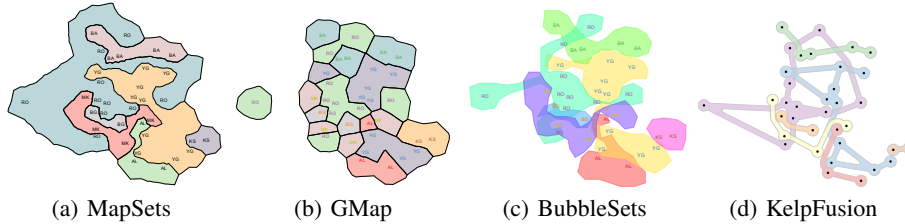
Here we compare our new algorithm MapSets with the existing approaches for map-like visualizations: GMap [11], BubbleSets [7], and KelpFusion [16]. A fully functional implementation of MapSets, GMap, and BubbleSets is available in an online system at <http://gmap.cs.arizona.edu>; see more details in Appendix B. The drawings of KelpFusion are courtesy of the authors [16].

Our first example is the senator voting graph; see Fig. 6. The vertices in the graph are the U.S. senators in 2010 positioned according to their home-cities in the U.S. The clustering is based on the political party they represent, red for republicans and blue for democrats. Clearly, both clustering and geographic information of the vertices are fixed and cannot be changed. One can see that GMap produces fragmented clusters, while BubbleSets and KelpFusion compute overlapping regions. On the other hand, the result of MapSets is contiguous and non-overlapping, which makes it easier to analyze the distribution of senators over the map.

The second example shows the population structure within Europe [18]. The original points correspond to genetic data from 1,387 Europeans (but we sampled only 50 vertices corresponding to Eastern Europe for illustration purposes). The positions of the



**Fig. 6:** The senator voting graph (the part of the U.S. west of Mississippi). The vertices are senators (red republicans and blue democrats) positioned according to their home-cities.



**Fig. 7:** The graph of genetic similarities between 50 individuals in Europe. The layout is computed using the principal component analysis, while the clusters correspond to the countries of origin of the individuals.

vertices come from the original principal component analysis, based on the similarity matrix. As the authors point out, the PCA plot (appropriately rotated) closely matches the geographic outlines of Europe; hence, it is undesirable to change the node positions. The clusters are extracted independently and corresponds to the countries of origin of the individuals. Again, only MapSets constructs non-fragmented disjoint regions; see Fig. 7. Arguably, this is easier to analyze than the overlapping regions produced by BubbleSets and KelpFusion; see more examples in Appendix A.

We next analyze the performance of our ink minimization heuristic. To this end, we utilize a collection of 9 real-world networks, that are embedded and clustered using the GMap tool with the default setting [11]. Table 1 gives details about the graphs and measurements of our ink saving algorithm; see Appendix B for more details on the datasets. Here,  $ALG$  shows the ratio of the total ink of the computed trees to the total length of the minimum spanning trees computed individually for every cluster. In other words, this is an approximation factor achieved by our algorithm on the test cases. Although we can only guarantee factor  $k\rho$ , in practice the algorithm performs very well, always producing a solution at most 1.6 times worse than the optimal. Our experiments indicate that ink minimization strategy often results in aesthetically more pleasant map visualizations; see Fig. 9 (in Appendix) for a comparison.

Similarly,  $ALG_{fd}$  indicates the utilized ink after the force-directed adjustments. As expected, the ink increases after the step, but the increase is not significant. On the other hand, the adjustments improve the quality of the resulting regions. Fig. 10 (in Appendix) provides an example of the maps computed with and without the heuristic; notice the narrow channels in the figure computed without the adjustments.

| graph        | $ P $ | $k$ | ALG   | $ALG_{fd}$ |
|--------------|-------|-----|-------|------------|
| Colors       | 50    | 6   | 1.002 | 1.012      |
| GD           | 506   | 23  | 1.582 | 1.612      |
| Recipes      | 381   | 15  | 1.356 | 1.502      |
| Trade        | 211   | 8   | 1.101 | 1.259      |
| Universities | 161   | 8   | 1.366 | 1.443      |
| SODA         | 316   | 11  | 1.204 | 1.296      |
| IPL          | 336   | 11  | 1.337 | 1.414      |
| SOCG         | 500   | 11  | 1.492 | 1.601      |
| TARJAN       | 252   | 16  | 1.150 | 1.197      |
| ALGO         | 500   | 5   | 1.547 | 1.650      |

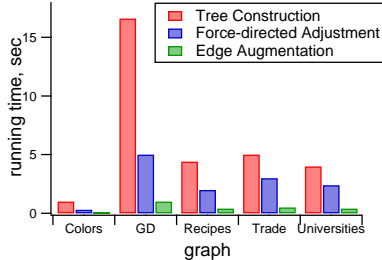
**Table 1:** Measurements of MapSets on test cases: ALG and  $ALG_{fd}$  stand for the ratio between the total ink of the drawing and the total length of the minimum spanning trees after the steps *Tree Construction* and *Force-directed Adjustment*, respectively.

We use a machine with Intel i5 3.2GHz and 8GB RAM for measuring running time; see Fig. 8. The algorithm is implemented in C++. Note that the last two steps, *Adding Auxiliary Points* and *Computing Regions*, are very efficient taking few milliseconds for the largest graphs, and hence are not included in the chart. The first step, *Tree Construction*, is usually the most time consuming; it is more efficient for nearly contiguous clusters (e.g., Colors) and less efficient for graphs with many fragments (e.g., GD). Although *Edge Augmentation* theoretically has cubic time complexity, it among the fastest steps in practice, because there are usually not many edges added. Overall, our algorithm processed all graphs (most with hundreds of vertices) in less than a minute. This is slower than the GMap and LineSets but comparable to BubbleSets. Since our algorithm extensively utilizes many primitive geometric operations (e.g., testing for segment intersections), using a specialized geometric library will likely improve the performance.

## 6 Conclusion and Future Work

We designed and implemented a new approach for visualizing embedded and clustered graphs. Unlike existing techniques, our MapSets method always produces contiguous and non-overlapping regions. Results of the initial evaluations seem promising. We also presented a simple approximation algorithm for the geometric problem of ink minimization motivated by the method. A natural future direction is to improve the approximation factor. It would be also worthwhile to carefully evaluate different convexity measures and select one that offers the best balance between ease of computation and visual quality of the resulting regions.

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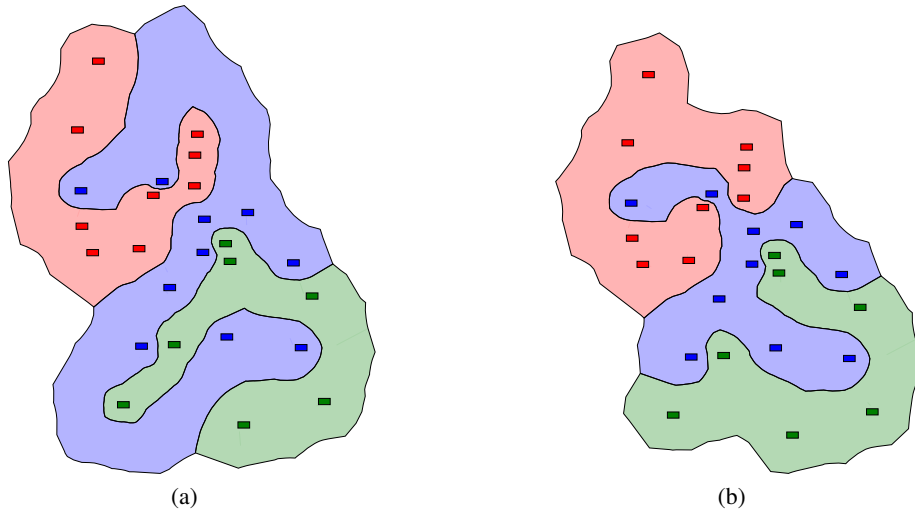


**Fig. 8:** Running times of the different steps of MapSets on some of the test cases.

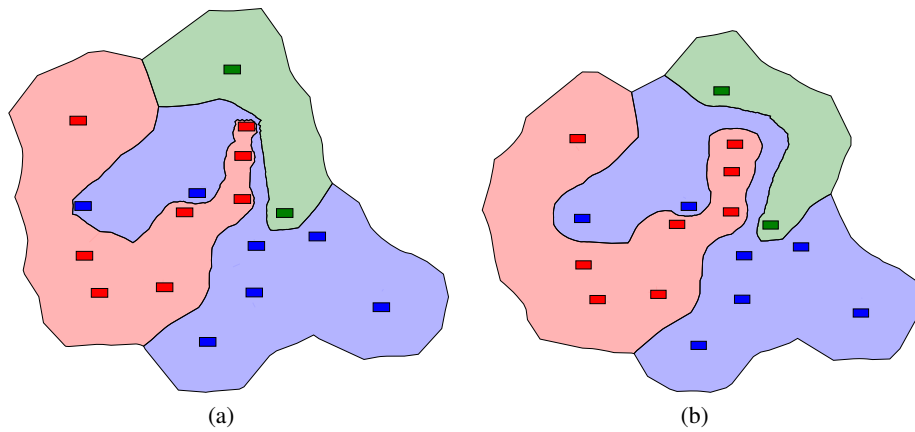
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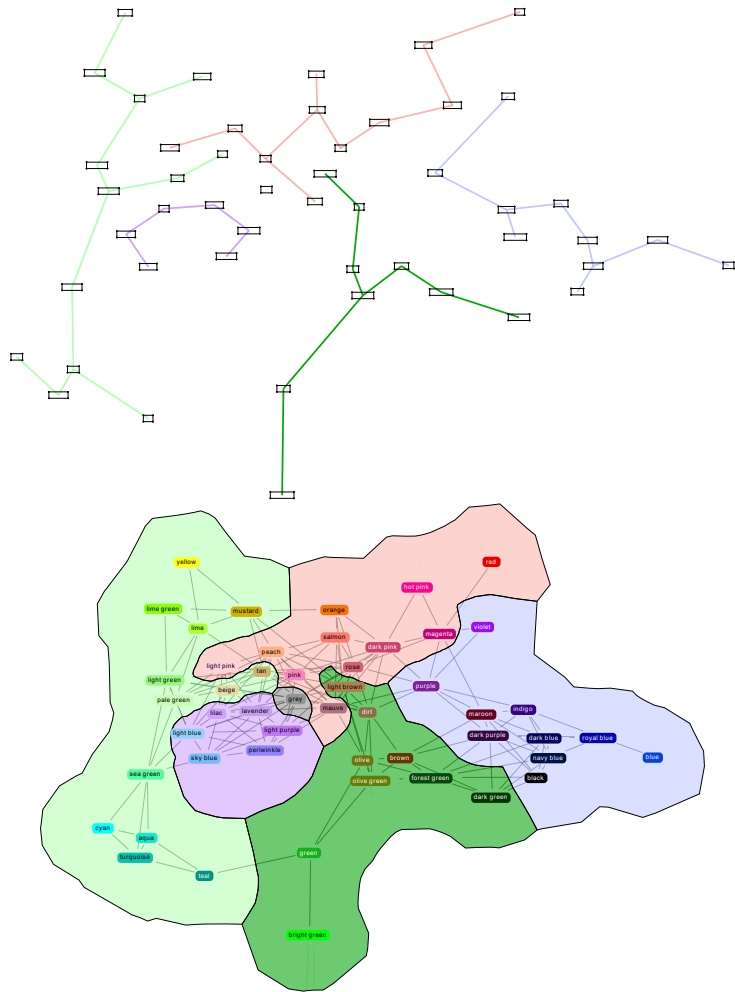
## A Appendix: Additional Figures



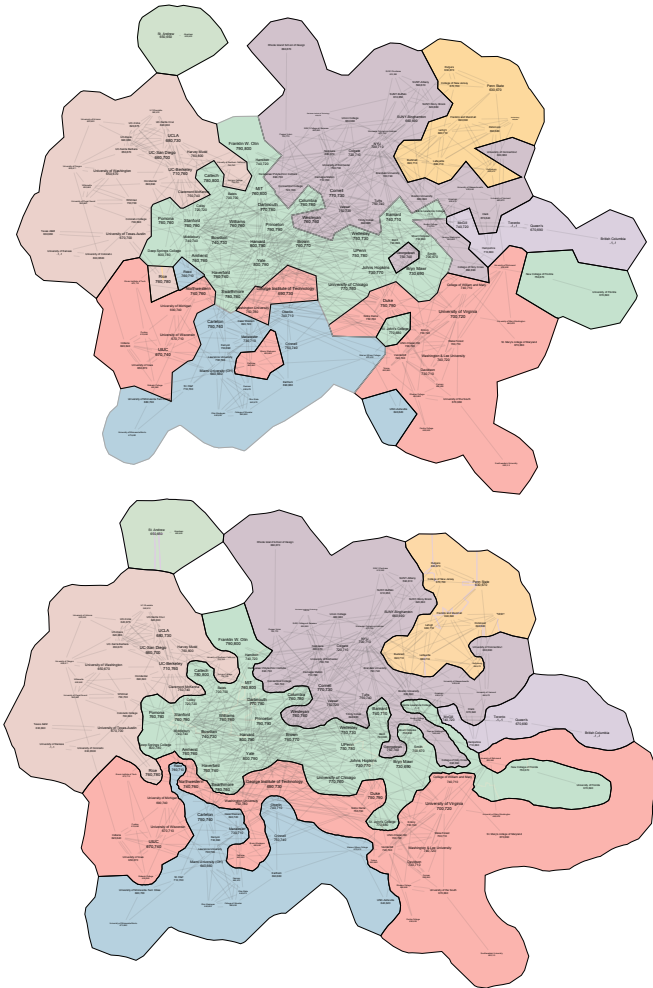
**Fig. 9:** The effect of ink minimization in MapSets. The results computed for the trees of total ink: (a) 6658 units ( $ALG = 1.06$ ) and (b) 6021 units ( $ALG = 1.172$ ).



**Fig. 10:** The effect of the *Force-directed Adjustment* step in MapSets: (a) without the adjustments and (b) with the adjustments. The very thin connection between two blue components is almost invisible without the force-directed adjustments.



**Fig. 11:** The graph is constructed using the 50 most common monitor colors. The edge-weights are defined by the distance in the RGB space between corresponding pairs. The layout is created using multidimensional scaling, clustering is done using the modularity optimization algorithm. (a) Computed non-intersected trees. (b) The final MapSets results. The image is zoomable and has high resolution.



**Fig. 12:** The Universities graph: network of the U.S. universities and their average SAT scores. The vertices are universities and edges are constructed based on similarities in admissions. The layout is computed using the force-directed algorithm, clustering is constructed by modularity optimization. (a) Fragmented map computed by GMap. (b) The contiguous MapSets result. The image is zoomable and has high resolution.

## B Appendix: Online System

A fully functional implementation of MapSets, GMap, and BubbleSets is available in an online system at <http://gmap.cs.arizona.edu>. In order to draw a graph, paste the graph in the *dot* format, press “Show Advanced Options”, then choose an appropriate visualization type and clustering/layout algorithm, and press “Create Map”. We implemented and made available the following visualization techniques:

- *Node-Link Diagrams* – the classical method for visualizing graphs;
- *GMap* – the Graph-to-Map algorithm for creating map-like visualizations [11];
- *BubbleSets* – the method suggested by Collins et al. [7], which is based on isocontours to overlay an arrangement of objects with enclosing set regions;
- *LineSets* – the set visualization technique by Alper et al. [2], that uses a simple curve traversing all of the elements of each set;
- *MapSets* – our new algorithm for visualizing embedded clustered graph.

We provide several classical graph layout algorithms (*force-directed algorithm*, *multidimensional scaling*) and clustering algorithms (*modularity clustering*, *K-means*, *hierarchical clustering*). The system supports the semantic zoom feature and interactive browsing, once can also save the image in a variety of formats; see Fig. 13. Source code for the entire system and all algorithms is available on GitHub.

**Graph Dataset** Real-world graphs used in our experiments can be downloaded from <http://gmap.cs.arizona.edu/datasets>. **ALGO**, **IPL**, **SOCG**, **SODA**, and **TARJAN** describe topics of research papers and contain the prominent words and phrases extracted from the titles of the papers. The edges represent similarities between the topics computed based on their co-occurrence in titles. **GD** is the co-authorship graph for the Graph Drawing Symposium; the vertices represent the authors and the edge represent papers published together. **Recipes** contain ingredients extracted from cooking recipes. **Trade** describes trade relationships between countries. The **Universities** dataset is based on average SAT scores in the U.S. universities.

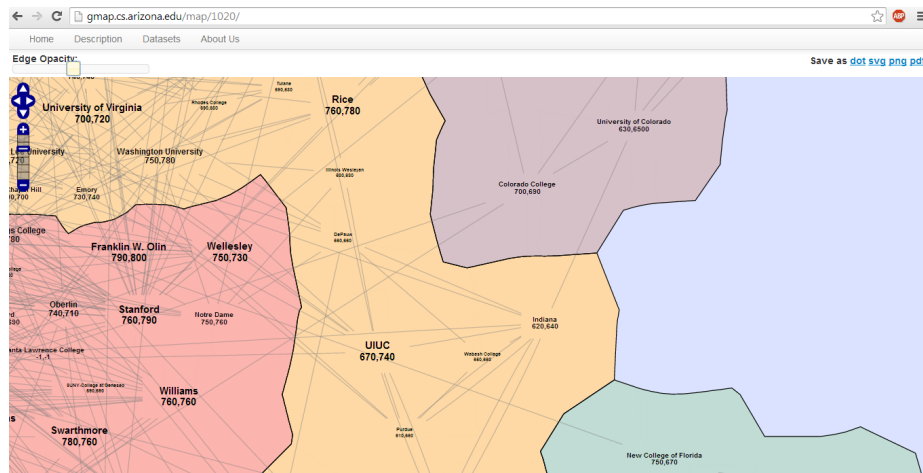


Fig. 13: The interface of our interactive system for visualizing clustered graphs.