# CLUSTERING PROPENSITY: A MATHEMATICAL FRAMEWORK FOR MEASURING SEGREGATION

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ABSTRACT. We propose a new family of metrics called CAPY (or *clustering propensity*) scores, designed to measure the clustering level of one or more subgroups within a population. The intended application is to offer new ways of measuring the segregation of demographic subgroups. We discuss two main capy scores, Edge and HalfEdge (as well as weighted variants of each) and we compare them to existing segregation scores in the political science, geography, and network science literature. To evaluate the scores, we compute and plot values of minority proportion  $\rho$  vs. clustering score C for test distributions on large  $n \times n$  grids, and on actual demographic data from U.S. states and cities. We argue that capy scores successfully discern qualitatively important differences while providing a stabler baseline for interpretation than classic scores like the Dissimilarity Index and Moran's I.

Keywords: Segregation, network clustering, assortativity, dissimilarity.

## **CONTENTS**



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#### 1. INTRODUCTION

<span id="page-1-1"></span><span id="page-1-0"></span>1.1. Background and goals. In this paper we present a family of "clustering propensity" scores that in part unites and in part adds to segregation and assortativity scores that already exist in the geography and network science literature. The goal is to present numerical tools for describing aspects of spatial distribution of populations that can help inform policy considerations.

We have set up the problem as follows: given a region of interest that has been partitioned into geographic units (such as census tracts or precincts), we construct a dual graph that records the geographic and demographic information. These dual graphs are flexible network structures that allow for mathematical analysis of spatial population distributions, which in turn leads to a very general framework for measuring segregation.

To analyze performance, we will consider a suite of questions aimed at evaluating whether a proposed score has adequate discernment and stability. That is, scores should offer a stable numerical baseline: similar scores should mean something qualitatively similar across scenarios; in particular, scores should not be heavily or chiefly sensitive to a non-pattern-related variable like city size, minority share, or choice of units. The units issue—in which changing the aggregation level has a drastic impact on output—is well known in the geography literature as a MAUP, or Modifiable Areal Unit Problem. Avoiding undue sensitivity to factors that are in some sense orthogonal to clustering or segregation will give us grounds to prefer capy scores to some classical alternatives. And at the same time, we will prefer scores that register meaningful qualitative differences in segregation scenarios.

This direction of investigation was motivated by the study of electoral redistricting. Demographic clustering has a major impact on political representation under the system of single-member districts that dominates the United States electoral scene. This is even made explicit in the checklist of features that must be established to bring a lawsuit under the Voting Rights Act of 1965—litigants must demonstrate that a minority group is "sufficiently large and geographically compact to constitute a majority in a single-member district" in order to press a claim that the group has been denied rightful representation.<sup>[1](#page-1-4)</sup> This phrasing acknowledges legally what is mathematically clear: the size of a minority population alone, without sufficient spatial clustering or "compactness," is not enough to guarantee that the group can secure representation in a districted system. We were motivated by wanting to measure clustering with tools compatible with statistical physics models, like the Ising model, that would allow us to design dynamical systems to intensify and relax the level of clustering and study the representational consequences. The intimate relationship between segregation and district-based representation will be discussed in future work.

#### 2. The theoretical framework of capy scores

<span id="page-1-3"></span><span id="page-1-2"></span>2.1. Geographical units and dual graphs. We begin by setting up definitions and notation to treat a city, state, or any other jurisdiction as a graph decorated with relevant demographic data. In our examples, we will use geographical units from the census, such as census tracts or census blocks, that partition the jurisdiction into pieces. The dual graph of a geographical partition is the graph formed by using a vertex (or node) to represent each unit, then connecting two vertices by an edge if the geographical units are adjacent. We can either adopt edges for *rook adjacency* (in which the shared boundary has to have positive length) or queen adjacency (in which we count units as being adjacent even if they just meet at a point). This is illustrated below in Figure [1.](#page-2-1)

At each node we can record demographic information for the geographic unit, including the total population and racial breakdown, based for instance on census data. The geographical units that make up a jurisdiction have populations of different sizes and compositions. Suppose we have two

<span id="page-1-4"></span><sup>&</sup>lt;sup>1</sup>In the VRA literature, this is called the Gingles 1 test. See Thornburg v. Gingles, 478 U.S. at 50, 1986.



<span id="page-2-1"></span>FIGURE 1. On the left is a partition of a region into five units. The middle and righthand figures represent dual graphs of this partition, where the middle figure has used rook adjacency and the righthand figure uses queen adjacency.

types of population, X and Y, such as Black and White residents.<sup>[2](#page-2-2)</sup> If the nodes of the dual graph are denoted  $v_i$ , then we can record integer-valued populations  $x_i$  and  $y_i$  in each unit, with total population  $p_i$  at the *i*th node. We may have  $p_i = x_i + y_i$  if each population member is classified in group X or group Y, or there may be other groups in the population. We will record the X population data as a vector  $\mathbf{x}: V \to \mathbb{Z}$ , and likewise write y for the Y population figures. For example, Figure [6](#page-10-0) shows the dual graph of the 99 counties in Iowa. The sizes of the nodes in the figure reflect 2010 Census population of the counties, which in fact varies by more than two orders of magnitude, from a minimum of 4029 to a maximum of 430,640.

The total population of a jurisdiction will be denoted  $\bar{p} = \sum_i p_i$ , and likewise  $\bar{x}$  and  $\bar{y}$  represent the total number of residents of X or Y type, respectively. We will introduce the notation  $\rho = \bar{x}/\bar{p}$ to represent the proportion of population X in the population at large, so that  $0 \leq \rho \leq 1$ . Since we typically focus on a population in the numerical minority, most of the plots will have  $0 \leq \rho < 1/2$ .

<span id="page-2-0"></span>2.2. The exploded graph and an inner product expression. We would like to measure the extent to which people of population X tend to live next to other people of population of X, rather than next to people of population Y. So we will classify within-unit adjacencies as well as adjacencies between neighboring units. There are scores for this in the literature when each node corresponds to a single person, but we have not found existing segregation scores that handle arbitrary percentages at each node of a network.

In the network science and applied mathematics literature, authors sometimes consider constructions that *aggregate* and *disaggregate* nodes in graphs; that is, a graph can be modified by collapsing a subgraph to a node, or by replacing a node with an appropriate subgraph. We will describe a massively disaggregated secondary graph associated to our dual graph which we call the *exploded graph*. We expand each node  $v_i$  into a complete graph (or *clique*)  $K_{p_i}$  on  $p_i$  nodes such that exactly  $x_i$  are of X type. If two nodes  $v_i$  and  $v_j$  are adjacent in the initial dual graph, then the exploded graph contains  $p_i \cdot p_j$  edges between the members of the respective cliques. This graph has an enormous number of nodes (one for each person in the jurisdiction) and edges, but it is a theoretical construction that we use to explain the logic of the main definitions; we note that the exploded graph never has to be built or stored.

We can define two expressions as follows:

$$
\langle \mathbf{x}, \mathbf{y} \rangle := \sum_{i} x_i y_i + \sum_{i \sim j} x_i y_j + x_j y_i ;
$$

$$
\langle \langle \mathbf{x}, \mathbf{y} \rangle \rangle := \frac{1}{2} \left( \sum_{i} \left( x_i y_i - \frac{x_i + y_i}{2} \right) + \sum_{i \sim j} x_i y_j + x_j y_i \right).
$$

<span id="page-2-2"></span><sup>2</sup>We note that census data includes a count of Black-only population and White non-Hispanic population, among many other racial classifications, including membership in more than one racial group. Census classification allows researchers to treat racial categories as though they are much more stable and clear than the social reality.



FIGURE 2. This figure shows the *exploded graph* associated to an initial graph with  $x = (4, 2)$ ,  $y = (3, 2)$ , and no other type of population. Here, the exploded graph has  $\langle x, y \rangle = 30$  edges between different-colored nodes,  $\langle x, x \rangle = 15$  edges between X nodes, and  $\langle \langle y, y \rangle \rangle = 10$  edges between Y nodes, making 55 edges in all. The proportion of X population in the jurisdiction is  $\rho = 6/11$ .

Here in both expressions the first summation is over all the nodes, and the second is over pairs of adjacent nodes. Note that the number of edges between populations X and Y within the clique associated to vertex *i* is  $x_i y_i$ , which means that  $\langle \mathbf{x}, \mathbf{y} \rangle$  is a precise count of the edges of XY type when X and Y are disjoint populations. On the other hand, the number of edges between two people of population X is

$$
\binom{x_i}{2} = \frac{x_i^2 - x_i}{2},
$$

so  $\langle x, x \rangle$  simplifies to a precise count of the number of edges of XX type.

We note another relationship between these expressions. Since quadratic terms dominate linear terms when the  $x_i$  and  $y_i$  are large, we get  $\langle x, y \rangle \approx 2 \langle\langle x, y \rangle\rangle$  for large populations.

Observe that  $\langle x, y \rangle$  is an inner product, so it has a nice representation in terms of matrix multiplication. Letting A be the adjacency matrix of the dual graph, we have

$$
\langle \mathbf{x}, \mathbf{y} \rangle = \mathbf{x}^T (A + I) \mathbf{y}.
$$

<span id="page-3-0"></span>2.3. Measuring clustering propensity. With the information above, we can define clustering propensity scores on the exploded graphs which have a clear probabilistic interpretation.

We can use this to define a one-sided score of the skew via

$$
\mathsf{Skew}(\mathbf{x}, \mathbf{y}) := \frac{\langle \mathbf{x}, \mathbf{x} \rangle}{\langle \mathbf{x}, \mathbf{x} \rangle + 2\langle \mathbf{x}, \mathbf{y} \rangle} = \frac{\langle \mathbf{x}, \mathbf{x} \rangle}{\langle \mathbf{x}, \ \mathbf{x} + 2\mathbf{y} \rangle}.
$$

Using the fact that  $\langle \mathbf{x}, \mathbf{y} \rangle \approx 2 \langle \langle \mathbf{x}, \mathbf{y} \rangle$ , we see that the skew is approximately  $\frac{\langle \langle \mathbf{x}, \mathbf{x} \rangle \rangle}{\langle \langle \mathbf{x}, \mathbf{x} \rangle + \langle \mathbf{x}, \mathbf{y} \rangle}$ , which is the ratio of the number of XX edges to the number of edges of either XX or XY type. In other words, among the edges that connect X population to either X or Y population, it records the share of XX edges. This measures the prevalence of X living next to X rather than Y, weighted by edges.

Therefore to devise a score of the clustering propensity between populations X and Y from an edge point of view, we can average the X and Y skews, arriving at the edge capy score

(1) 
$$
\mathsf{Edge}(\mathbf{x}, \mathbf{y}) := \frac{1}{2} \left( \frac{\langle \mathbf{x}, \mathbf{x} \rangle}{\langle \mathbf{x}, \mathbf{x} \rangle + 2 \langle \mathbf{x}, \mathbf{y} \rangle} + \frac{\langle \mathbf{y}, \mathbf{y} \rangle}{\langle \mathbf{y}, \mathbf{y} \rangle + 2 \langle \mathbf{x}, \mathbf{y} \rangle} \right)
$$

Note that the score can be extended to compare the clustering of multiple disjoint sets, such as with

$$
\mathsf{Edge}(\mathbf{x}, \mathbf{y}, \mathbf{z}) = \frac{1}{3} \left( \frac{\langle \mathbf{x}, \mathbf{x} \rangle}{\langle \mathbf{x}, \mathbf{x} \rangle + 2\langle \mathbf{x}, \mathbf{y} \rangle + 2\langle \mathbf{x}, \mathbf{z} \rangle} + \frac{\langle \mathbf{y}, \mathbf{y} \rangle}{\langle \mathbf{y}, \mathbf{y} \rangle + 2\langle \mathbf{x}, \mathbf{y} \rangle + 2\langle \mathbf{y}, \mathbf{z} \rangle} + \frac{\langle \mathbf{z}, \mathbf{z} \rangle}{\langle \mathbf{z}, \mathbf{y} \rangle + 2\langle \mathbf{x}, \mathbf{z} \rangle + 2\langle \mathbf{y}, \mathbf{z} \rangle} \right),
$$

and so on to arbitrarily many populations.

However, if we want to reframe this as a propensity in terms of the vertices (the people) rather than the edges (the adjacencies of people), it is more natural to set up the ratio in terms of half-edges rather than edges. A half-edge is a vertex-edge pair  $(v, e)$  in which edge e is incident to vertex v. The share of X type half-edges which belong to an XX edge is  $\frac{2\langle\langle \mathbf{x}, \mathbf{x}\rangle\rangle}{2\langle\langle \mathbf{x}, \mathbf{x}\rangle\rangle + \langle \mathbf{x}, \mathbf{y}\rangle}$ , which is asymptotic to

$$
\mathsf{Skew}'(\mathbf{x}, \mathbf{y}) = \frac{\langle \mathbf{x}, \mathbf{x} \rangle}{\langle \mathbf{x}, \mathbf{x} \rangle + \langle \mathbf{x}, \mathbf{y} \rangle}.
$$

This has the intuitively appealing interpretation as the probability that a neighbor of an X person is another X person rather than a Y person. Accordingly, we define the half-edge capy score to be

(2) HalfEdge(**x**, **y**) := 
$$
\frac{1}{2} \left( \frac{\langle \mathbf{x}, \mathbf{x} \rangle}{\langle \mathbf{x}, \mathbf{x} \rangle + \langle \mathbf{x}, \mathbf{y} \rangle} + \frac{\langle \mathbf{y}, \mathbf{y} \rangle}{\langle \mathbf{y}, \mathbf{y} \rangle + \langle \mathbf{x}, \mathbf{y} \rangle} \right),
$$

noting that it can just as easily be extended to more than two populations.

This will be the clustering propensity score that receives our strongest focus in this paper: it averages the average tendency of each subgroup of population to have members of their own subgroup, and not the other, as neighbors.

<span id="page-4-0"></span>2.4. Within-unit and between-unit weighting. A natural variant on these scores is to weight the connections within geographical units differently than those between neighboring units. To accomplish this, we choose  $\lambda \geq 0$  and set

$$
\langle \mathbf{x}, \mathbf{y} \rangle_{\lambda} := \lambda \left( \sum_{i} x_i y_i \right) + \sum_{i \sim j} x_i y_j + x_j y_i.
$$

With this, we can simply repeat the formulas for clustering scores using the weighted inner products, such as

$$
\mathsf{HalfEdge}_\lambda(\mathbf{x}, \mathbf{y}) := \frac{1}{2} \left( \frac{\langle \mathbf{x}, \mathbf{x} \rangle_\lambda}{\langle \mathbf{x}, \mathbf{x} \rangle_\lambda + \langle \mathbf{x}, \mathbf{y} \rangle_\lambda} + \frac{\langle \mathbf{y}, \mathbf{y} \rangle_\lambda}{\langle \mathbf{y}, \mathbf{y} \rangle_\lambda + \langle \mathbf{x}, \mathbf{y} \rangle_\lambda} \right).
$$

In this way, any normalization factor one might introduce for  $\langle , \rangle_{\lambda}$  cancels out of the numerator and denominator, and we obtain a score that weights the two kinds of neighbors differently.

For instance, if one is working with geographical units that are chosen in part for their social unity, such as census tracts, then it would be reasonable to weigh the within-tract adjacencies more heavily than those between neighboring tracts, such as by taking  $\lambda = 2$  or  $\lambda = 5$ . If the units are counties, then there are some states in which people identify strongly with their county, such as Texas, and other states in which most people don't know what county they live in, such as Massachusetts. Some choice of  $\lambda$ -weighting could then be appropriate for studies of changing segregation over time in Texas.

Note that as  $\lambda \to \infty$ , the vertex terms dominate the weighted terms, so that in the limit we have  $\lim_{\lambda\to\infty}\langle x,y\rangle_{\lambda}=\sum_{i}x_{i}y_{i}$ . This defines the following weighted CAPY scores in the limit, defined by summing over the geographical units.

$$
\mathsf{HalfEdge}_{\infty}(\mathbf{x}, \mathbf{y}) = \frac{1}{2} \left( \frac{\sum x_i^2}{\sum x_i (1 + x_i y_i)} + \frac{\sum y_i^2}{\sum y_i (1 + x_i y_i)} \right).
$$

Of course, because the interaction between neighboring nodes has been dropped out, this becomes a node-based score (i.e., ignoring edges) like several classical scores discussed in the next section  $(S3.1)$ .

## 3. Comparison to existing literature

<span id="page-4-1"></span>We will survey some of the numerous existing segregation scores in the social science and applied mathematics literature, translating them into the notation of this paper for ease of comparison. Recall that **p** is the vector of population at each node, and  $\bar{x}, \bar{y}, \bar{p}$  are the jurisdiction-wide populations of X type, Y type, and all residents, respectively. We also have  $\rho$  as the jurisdiction-wide proportion of X population, and  $\rho_i = x_i/p_i$  the proportion at node i.

<span id="page-5-0"></span>3.1. Node-based scores: Dissimilarity, Frey, and Gini. The segregation literature has three major scores that have been described as measuring "evenness," or the consistency of the levels of a sub-population over the units that make up a jurisdiction.

$$
D(\mathbf{x}) = \frac{1}{2\bar{x}(\bar{p} - \bar{x})} \sum_{i} |x_i \bar{p} - p_i \bar{x}| \; ; \qquad F(\mathbf{x}, \mathbf{y}) = \frac{1}{2\bar{x}\bar{y}} \sum_{i} |x_i \bar{y} - y_i \bar{x}| \; ;
$$

$$
G(\mathbf{x}) = \frac{1}{2\bar{x}(\bar{p} - \bar{x})} \sum_{i,j} |x_i p_j - p_i x_j|.
$$

These are called the Dissimilarity score, the Frey index, and the segregation Gini index, respectively. We note that all three are based on a similar determinant-like expression:  $|vw'-wv'|$  can be interpreted as twice the area of the triangle described by vector  $(v, w)$  and vector  $(v', w')$ , as in Figure [3.](#page-5-2)



<span id="page-5-2"></span>Figure 3. This area term is only zero if the vectors point the same direction, which occurs when there is an equality of ratios:  $\frac{v}{w} = \frac{v'}{w'}$ .

So all three of these formulas, while set up slightly differently from one another, measure how even the distribution of population X is:

- $D(\mathbf{x})$  measures how closely the unit proportions  $\rho_i = \frac{x_i}{p_i}$  $\frac{x_i}{p_i}$  line up with the citywide proportion  $\rho = \frac{\bar{x}}{\bar{x}}$  $\frac{x}{\bar{p}};$
- $F(\mathbf{x}, \mathbf{y})$  measures how nearly two groups X and Y have equal proportion of each unit's population;
- $G(\mathbf{x})$  looks over all pairs of units and measures how nearly  $\rho_i = \rho_j$ .

The determinental interpretation of the scores makes it easy to see that  $D(\mathbf{x}) = F(\mathbf{x}, \mathbf{p} - \mathbf{x})$ , so Frey's index can be seen as a generalization of dissimilarity to pairs of (not necessarily complementary) populations.[3](#page-5-3)

Dissimilarity and this Gini score (which borrows its name from the more famous area-based index of wealth distribution) are among the 20 segregation scores discussed in the classic Massey–Denton survey of segregation indices [\[8\]](#page-11-1). This or very similar formulations of Dissimilarity go back to at least the 1950s and have been much used and discussed since then (see [\[2,](#page-11-2) [5,](#page-11-3) [8\]](#page-11-1) and their references).

Note that each of these three scores is given by summing over the nodes without reference to adjacency, none of them can take into account the spatial relationship between geographic units, so they all treat neighboring units no differently than units on opposite sides of a city.

<span id="page-5-1"></span>3.2. Spatial scores in the geography literature, including Moran's I. Many authors in the geography literature have attempted to modify these scores to take spatial relationships between units into account by "spatial weighting," which can be set up to take into account when units are adjacent, or within a fixed distance, or simply to upweight pairs of units when they are relatively closer or share longer boundary segments. For instance Dawkins in two papers in the 2000s [\[3,](#page-11-4) [4\]](#page-11-5) provides spatialized variants of the Gini score from the last section.

<span id="page-5-3"></span><sup>&</sup>lt;sup>3</sup>In the papers of Frey, the index we call F is referred to as dissimilarity and denoted D, for example in [\[7\]](#page-11-6).

But the most widely used spatial statistic is very likely Moran's I, introduced in 1950 by a statistician named P.A.P. Moran. Consider a node-wise value  $\mathbf{x} = (x_1, \ldots, x_n)$ , such as population of group X in our setup. Let  $x_0 = \bar{x}/n$  be the average level over the nodes. We might choose to translate **x** so that its mean is zero, defining  $\mathbf{v} = (x_1 - x_0, \dots, x_n - x_0)$ . Then we can define

$$
I = \frac{n}{|E|} \cdot \frac{\sum_{i \sim j} (x_i - x_0)(x_j - x_0)}{\sum_i (x_i - x_0)^2} = \frac{n}{|E|} \cdot \frac{\mathbf{v}^T A \mathbf{v}}{\mathbf{v}^T \mathbf{v}}
$$

,

in terms of the adjacency matrix  $A$ , which in linear algebra terms is just a normalized Rayleigh quotient for the vector v.

To compute this for several test patterns, notice that it can be interpreted as the average of  $v_i v_j$ values for adjacent pairs of units divided by the average  $v_i^2$  over the single units. Moran's coefficient for a checkerboard pattern of 0 and 1 on a grid graph would be  $-1$ , because every  $v_i = x_i - x_0$ would be  $\pm 1/2$ , but all of the signs in the numerator would be negative because of the alternation. On the other hand, uniformly distributing 0 and 1 values on the vertices of a large graph would give a score near  $I = 0$ , because of the expected cancellation of positive (like) and negative (unlike) terms. And a heavily clustered 0-1 configuration would tend toward  $I = 1$ , because nearly all  $v_i v_j$ terms would be between like pairs, giving  $v_i v_j = v_i^2$ , and the two types of adjacency occur in the same proportion as the two types of nodes.

A local version of this score has been proposed, defined in the neighborhood of the jth unit. This can be useful to locate clustering. It can be defined by

$$
I_j = n(x_j - x_0) \cdot \frac{\sum_{i \sim j} (x_i - x_0)}{\sum_i (x_i - x_0)^2},
$$

which is just like the global  $I$  except that the numerator only looks at adjacencies involving node  $j$  and we have dropped the normalization by the total number of edges. This has been applied to redistricting in work of Chen–Rodden [\[1\]](#page-11-7).

One important critique of Moran's I is that it is heavily subject to MAUP, or the modifiable areal unit problem discussed in the introduction. This is an important concern in geography: if a score depends too heavily on the choice of geographical units—such as census blocks versus block groups, tracts, etc—that undermines its diagnostic usefulness. To see this problem in Moran's I, consider again the 0-1 checkerboard configuration on a large grid. If the individual units are used, we get  $I = -1$ , but if we reaggregate mildly so that the 2 × 2 pieces are used as units, then each unit has an identical composition and we get  $I = 0$ .

<span id="page-6-0"></span>3.3. Assortativity scores in network science. In network science, techniques from graph theory, geometry, and data analysis are used to study the structure of networks that come from real-world data. The field largely developed through applications to ecology, epidemiology, and social networks. The term *assortativity* is attached to a range of network scores that are broadly designed to assess whether nodes are more often adjacent to nodes like or unlike themselves, making it precisely aligned with the motivation used to define capy scores above. Some of the early focus in the study of assortativity was on graph-theoretic properties, asking for instance whether neighbors are likely to have similar degree or connectivity properties. But demographic sorting has also been considered. For instance, one common example is to study the racial assortativity of social networks; this is clearly relevant to the current application, which is racial assortativity of geographical networks. With an example like this in mind, a recent survey by Mark Newman [\[9\]](#page-11-8) gives as its main example an assortativity coefficient Q that had been developed to study the spread of HIV. Generally defined with respect to any number of non-overlapping groups that make up a population, it simplifies to something familiar in the case of a group and its complement: it is built from the fraction of XX edges among the XX and XY edges and the corresponding term for YY.

$$
Q=\left[\frac{\langle\!\langle \mathbf{x},\mathbf{x}\rangle\!\rangle}{\langle\!\langle \mathbf{x},\mathbf{x}\rangle\!\rangle+\langle \mathbf{x},\mathbf{y}\rangle}+\frac{\langle\!\langle \mathbf{y},\mathbf{y}\rangle\!\rangle}{\langle\!\langle \mathbf{y},\mathbf{y}\rangle\!\rangle+\langle \mathbf{x},\mathbf{y}\rangle}\right]-1.
$$

Dropping the linear terms (so that  $\langle x, x \rangle \approx 2 \langle x, x \rangle$ ), we have  $Q \approx 2$ Edge − 1, which means that it captures just the same information as Edge, but affinely rescaled to vary over [−1, 1] rather than  $[0, 1]$ .

Thus assortativity is in a sense already in the capy family. However, Q only handles nodes whose attributes vary over a finite set, and our exploded graph construction enables us to deal with percentage values, which is a significant generalization. In addition, we think that the HalfEdge score is a valuable variant on the edge-centered view.

### 4. Asymptotics on grid graphs

<span id="page-7-0"></span>We derive the theoretical behavior of the edge and half-edge CAPY scores in different configurations. Consider an  $n \times n$  grid with each node holding a population of M people, so that the total population of the grid is  $\bar{p} = Mn^2$ . We recall that  $\rho = \bar{x}/\bar{p}$  (so that  $0 \le \rho < 1/2$ ) is the parameter representing the (minority) proportion of population X in the grid. In this section we will analyze scores asymptotically as  $n \to \infty$ .

#### <span id="page-7-1"></span>4.1. Test configurations on asymptotic grids.

4.1.1. Perfect checkerboards. A perfect checkerboard configuration with density  $\rho$ , which we call Checkerboard and denote by  $Ch_{\rho}$ , alternates between  $x_i = 0$ ,  $y_i = M$  and  $x_j = 2\rho M$ ,  $y_j = (1-2\rho)M$ on adjacent nodes. In this way it maintains the global proportion  $\rho$  of population X.

That is, the pattern of population X is made up of repeating blocks of the form  $\begin{bmatrix} 2\rho & 0 \\ 0 & 2 \end{bmatrix}$  $0 \quad 2\rho$  . This gives

$$
\sum_{i=1}^{n} a_i
$$

$$
\langle \mathbf{x}, \mathbf{x} \rangle = \frac{n^2 4\rho^2 M^2}{2};
$$
  
\n
$$
\langle \mathbf{y}, \mathbf{y} \rangle = \frac{n^2 M^2 + n^2 M^2 (1 - 2\rho)^2}{2} + 4n^2 M^2 (1 - 2\rho);
$$
 and  
\n
$$
\langle \mathbf{x}, \mathbf{y} \rangle = \frac{n^2 M^2 2\rho (1 - 2\rho)}{2} + 2n^2 M^2 2\rho.
$$

The capy scores become

$$
\mathsf{Edge}(Ch_{\rho}) = \frac{25 - 50\rho + 20\rho^2 - 4\rho^3}{2(5 - \rho)(5 - 2\rho^2)} \quad \text{and} \quad \mathsf{HalfEdge}(Ch_{\rho}) = \frac{5 - 8\rho}{2(5 - 5\rho)}.
$$

4.1.2. Constant/uniform distributions. Next, consider the constant or uniform configuration  $Const_o$ , where each node has  $x_i = \rho M$  and  $y_i = (1 - \rho)M$ . Then,

$$
\langle \mathbf{x}, \mathbf{x} \rangle = n^2 M^2 \rho^2 + 2n^2 M^2 2\rho^2;
$$
  
\n
$$
\langle \mathbf{y}, \mathbf{y} \rangle = n^2 M^2 (1 - \rho)^2 + 2n^2 M^2 2 (1 - \rho)^2;
$$
 and  
\n
$$
\langle \mathbf{x}, \mathbf{y} \rangle = n^2 M^2 \rho (1 - \rho) + 2n^2 M^2 2\rho (1 - \rho).
$$

The capy scores are then

$$
\mathsf{Edge}(Const_{\rho}) = \frac{1 - \rho + \rho^2}{2 + \rho - \rho^2} \quad \text{and} \quad \mathsf{HalfEdge}(Const_{\rho}) = \frac{1}{2}.
$$

4.1.3. Isolated configurations. Next, consider binary grid configurations in which no two nodes with X population are adjacent. For a given  $\rho$ , there must be  $\rho n^2$  nodes of X type to get a total X proportion of  $\rho$ . Any such configuration is called an *isolated* configuration, and denoted Isol<sub> $\rho$ </sub>. We compute

$$
\langle \mathbf{x}, \mathbf{x} \rangle = n^2 M^2 \rho; \n\langle \mathbf{y}, \mathbf{y} \rangle = n^2 M^2 (1 - \rho) + 2(2n^2 - 4n^2 \rho) M^2; \text{ and } \n\langle \mathbf{y}, \mathbf{y} \rangle = 4n^2 M^2 \rho.
$$

We get

$$
\mathsf{Edge}(Isol_\rho) = \frac{25 - 41\rho}{9(5 - \rho)} \quad \text{and} \quad \mathsf{HalfEdge}(Isol_\rho) = \frac{3 - 5\rho}{5 - 5\rho}.
$$

4.1.4. Clusters. As in the isolated configuration, the one-cluster configurations  $OneClust_{\rho}$  will have  $x_i = 0$  or M at each node. But this time the  $\rho n^2$  nodes of type X are in a single large cluster. The only contributions to the count of XY edges  $(\langle x, y \rangle)$  will be the *perimeter* of the X cluster. We will choose the cluster to be a asymptotic to the square with side length  $\sqrt{\rho}n$ , giving  $2n^2\rho$  XX edges and  $2n^2(1-\rho)$  YY edges to first order, i.e., up to an error term that is linear rather than quadratic in  $n$ . We have

$$
\langle \mathbf{x}, \mathbf{x} \rangle = n^2 M^2 \rho + 4n^2 M^2 \rho; \n\langle \mathbf{y}, \mathbf{y} \rangle = n^2 M^2 (1 - \rho) + 4n^2 M^2 (1 - \rho);
$$
 and 
$$
\langle \mathbf{x}, \mathbf{y} \rangle = 2n M^2 \sqrt{\rho},
$$

with CAPY scores

$$
\mathsf{Edge}(OneClust_{\rho}) = \mathsf{HalfEdge}(OneClust_{\rho}) = 1.
$$

In Section [4.3,](#page-8-1) we will plot configurations with one and multiple clusters to illustrate how, as the perimeter of minority clusters increased, the capy scores decrease.

<span id="page-8-0"></span>4.2. Asymptotic comparisons. We can plot the four test configurations over  $0 < \rho < \frac{1}{2}$ .

<span id="page-8-1"></span>4.3. Corroboration on finite grids. To test our analysis of the capy scores for clustering, we generated test configurations as described in the last section on a  $90 \times 90$  grid graph, where each unit has a population of 1000. We plot the following configurations for  $\rho = .1, .2, .3, .4, .5$ .

- Isolated configurations where some cells are entirely X and no X cell has any rook-adjacent X neighbors;
- One cluster in which cells are entirely X;
- Two to ten clusters of cells that are entirely X;
- Checkerboard where cells alternate between  $x_i = 2\rho$  and 0;
- Constant X population of  $\rho$  in each cell.

The results, plotted in Figure [4,](#page-9-0) are nicely consistent with the theoretical behavior derived above, showing that the asymptotic calculations already work at that scale.

#### 5. Observed network example: Counties of Iowa

<span id="page-8-2"></span>Many papers in computational social science propose network-based scores but only try them on grid graphs. We next move to the real-data setting that is as close as possible to the grid configuration: the 99 counties of Iowa, whose (rook) dual graph is extremely patterned, with triangle and square structure. Besides being slightly more combinatorially complex than a grid, it also has substantial variation in the population by node, as noted above. We carry out HalfEdge calculations on the test configurations from above, as follows:



<span id="page-9-0"></span>FIGURE 4. CAPY scores for test patterns, on asymptotic grids (top) and  $90 \times 90$  grid graphs (bottom).



Figure 5. These are one-cluster, perfect checkerboard, and isolated configurations, respectively, on a large grid. Each has  $\rho = 40\%$  minority population.

- Isolated configurations are produced by randomly choosing nodes to fill with X population (shown in cyan) such that no two X nodes are adjacent;
- Constant configurations are produced by varying  $\rho$  from 0 to 1/2 and giving each node that share of X population;

• One-cluster configurations are produced by randomly choosing nodes to be all X and growing the cluster by adding random neighbors.

The results are shown in Figure [6.](#page-10-0) We do not feature checkerboard configurations, since those are only defined on bipartite graphs.



<span id="page-10-0"></span>Figure 6. The left-hand side shows one example each of an isolated, uniform, and one-cluster configuration of populations  $X$  (cyan) and  $Y$  (magenta) on the dual graph for Iowa's counties. By generating thousands of test configurations in these patterns at different levels of  $\rho$  (the proportion of population of X type), we can observe trends in the HalfEdge score.

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# Data Appendices

## Moon Duchin, Tyler Piazza

#### Appendix A. Description of data

<span id="page-12-0"></span>We have chosen 100 metropolitan areas (Metros) with significant geographical and demographic variation, including all of the top 40 most populous metro areas in the continental United States (by 2010 Census population).

We began with the Core Based Statistical Area (CBSA) shapefiles from the year 2013, fixing these as the definitions of the 100 Metros. We then intersected these with census tracts from 1990, 2000, and 2010, creating three timestamps with a constant geographical extension in which to compare change over time. Demographic data were joined to these tracts and partial tracts from NHGIS data by Python scripts acting on json files. (See GITHUB for all data and code.) Tracts with zero population were removed from the dataset, and the corresponding nodes removed from the dual graphs (though if two vertices were mutually adjacent to a zero-population tract, they were made adjacent to one another after deletion).

In this appendix, we present a range of tables and plots to illustrate features of segregation scores computed on these U.S. Metros. For consistency, we will fix two population subgroups to compare: White and POC. "White" denotes white non-Hispanic Census population, while "POC" (or people of color) represents the complement of this, encompassing all other racial and ethnic groups.

#### Appendix B. Comparisons and observations

<span id="page-12-1"></span>B.1. Change over time. We used the three timestamps in our dataset to construct plots to represent the change in segregation over time as reflected in these scores.



Figure 7. Capy scores for 5 large and 5 medium-sized Metros (population over 1.8 million and 1-1.8 million, respectively) at three timestamps. Most are getting more diverse and less segregated over time.



B.2. Comparing the scores pairwise. One notable feature of these scores is that they disagree significantly from one another on how to rank the 100 Metro areas.

Figure 8. Pairwise comparisons of how the segregation scores rank the 100 Metro areas with respect to 2010 data.

B.3. Within-tract and between-tract measurements. The next set of plots reports the differences that are imposed by varying the weighting of within-tract comparisons relative to betweentract comparisons. Recall that  $\lambda = \infty$  is the node-only variant (i.e., which disregards adjacency of tracts) and at the other extreme  $\lambda = 0$  is the edge-only variant.

We find that the Capy scores are making nontrivial use of the tract adjacency patterns, as reflected modest but visible change in rankings as  $\lambda \to \infty$ . Thus the score is actually and not just theoretically sensitive to the spatial arrangement of tracts. In the other direction, the finding is more surprising: varying  $0 \leq \lambda \leq 1$  has virtually no effect at all on the Metro rankings. This means that there is essentially no information loss in practice when discarding the within-tract scoring.



Figure 9. These plots compare the Capy scores to their weighted variants, where the edge terms are weighted  $\lambda$  times as heavily as the node terms.

B.4. Stability. Ideally, a segregation score should not simply reflect information that is more simply captured by an aggregate Metro statistic, such as the size of the city, the POC share of the population, or the choice of units. We address the choice of units below in [§B.6.](#page-16-0) Here we consider the relationship with city size and POC share.



FIGURE 10. These plots compare Capy scores to the POC Share  $(\rho)$  in and population of each Metro area. The last row has cities with population less than 1.8 million.

HEdge tends to score the whitest Metros at .5, which is the lowest score ever observed—so it's systematically scoring the whitest metro areas as the least segregated. The Edge score also tends to .5, but that's right in the middle of the observed scores.

B.5. Edge vs. half-edge discrepancy. Half-edge is better set up for earth-mover notion of segregation



Figure 11. Are the differences between scores primarily explained by factors orthogonal to segregation?



<span id="page-16-0"></span>B.6. Modifiable Areal Unit Problem. The MAUP.



Figure 12. There are about 3 times as many block groups as tracts in Chicago, and about 22 times as many blocks as block groups.



B.7. Tests on manufactured grids. Tests on generated grids.

Figure 13. Average values from tests where a 100 by 100 grid is filled in a checkerboard with alternating 1 plus epsilon or 0 plus epsilon. Epsilon is a uniform random variable in [−0.001, 0.001]. The 50 by 50 grid was made from the 100 by 100 grid by compressing 2 by 2 blocks into a single block.



Figure 14. All on 100 by 100 grids

Total population of each square is 1, possibly plus epsilon. Epsilon  $(\epsilon)$  is generally a uniform random variable in  $[-0.001, 0.001]$ . In the graph with "half of the board given positive  $\epsilon$ ", we specified that the epsilons on the left half of the grid were non negative, and the epsilons on the right half were non positive. For the one cluster tests, we created rectangles which were as close to a square as possible while having area equal to  $100.100 \cdot \rho$ . The width, height pairs of the clusters for the  $\rho$  values in increasing order are: (20, 25),(25, 40),(30, 50),(40, 50),(50, 50),(50, 60),(50, 70),(50, 80),(60, 75),(50, 100).

## Appendix C. Tabular results

<span id="page-19-0"></span>

CLUSTERING PROPENSITY 21



This table of 100 metropolitan areas has the scores for 3 decades (2010, 2000, 1990) for the scores Edge, Half Edge, Dissimilarity, and Moran's I (E,HE,D,I respectively).



CLUSTERING PROPENSITY 23



This table of 100 metropolitan areas has the ranks of the scores for 3 decades (2010, 2000, 1990) for the scores Edge, Half Edge, Dissimilarity, and Moran's I (E,HE,D,I respectively). A rank of 1 in  $E'10$  means that the score was the largest Edge score of the 100 metropolitan area scores for Edge of 2010.

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