

Article



Using Ramsey Theory to Measure Unavoidable Spurious Correlations in Big Data

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Abstract: Given a dataset, we quantify the size of patterns that must always exist in the dataset. This is done formally through the lens of Ramsey theory of graphs, and a quantitative bound known as Goodman's theorem. By combining statistical tools with Ramsey theory of graphs, we give a nuanced understanding of how far away a dataset is from correlated, and what qualifies as a meaningful pattern. This method is applicable to a wide range of datasets. As examples, we analyze two very different datasets. The first is a dataset of repeated voters (n = 435) in the 1984 US congress, and we quantify how homogeneous a subset of congressional voters is. We also measure how transitive a subset of voters is. Statistical Ramsey theory is also used with global economic trading data (n = 214) to provide evidence that global markets are quite transitive. While these datasets are small relative to Big Data, they illustrate the new applications we are proposing. We end with specific calls to strengthen the connections between Ramsey theory and statistical methods.

Keywords: statistics; data analysis; Ramsey theory; graph theory; transitivity

1. Introduction

In the realm of data science, the conventional wisdom is that "more data is always better", but is this the case? As a dataset *D* becomes larger, Ramsey theory describes the mathematical conditions by which disorder becomes impossible. The impossibility of disorder is analogous to the existence of unavoidable and spurious correlations in large datasets. This makes understanding and measuring the extent of these spurious correlations essential in any attempt to glean meaningful information about *D*. Ramsey theory is a productive area of mathematical research, and there are many types of systems with Ramsey results, as explored in 2016 by Calude and Longo [1]. There, the authors asked the question, how can Ramsey theory be used to understand spurious and unavoidable correlations in data science?

For example, the pigeonhole principle is an extreme, basic version of the Ramsey statement, "if a given person wears 8 different shirts in a given week and only has 7 shirts, then there must have been a single shirt that was worn at least twice". Here, the dataset is the collection of shirts, with each shirt assigned a day. The unavoidable spurious correlation is that (at least) one shirt is worn twice. It would be incorrect to conclude that the given person has a particular affinity for that repeated shirt. However, in this case, there is no meaningful conclusion we can draw, despite the natural human desire to attribute meaning to a pattern that is observed but forced to exist by the pigeonhole principle.

However, we might try to draw meaningful conclusions if we identify a shirt that the person wore three times in the same week because the pigeonhole principle on its own cannot guarantee these

beyond the base requirement that there is a single shirt that must be worn twice in a given week. This leads to our major connection between Ramsey theory and statistical analysis:

Remark 1 (Spurious Correlations through Ramsey theory). *Ramsey theory ensures that patterns and correlations must always exist in a sufficiently large system. Meaningful correlations are measurements in that system that are over and above the minimum amount guaranteed by Ramsey theory.*

Goodman's formula [2] provides a way to calculate the required number of certain relationships in a relational database. We use Goodman's formula to quantify how many correlations must be observed to ensure that some of the correlations are not spurious. Put another way, we use Goodman's formula to test the null hypothesis H_0 that a graph representing the relationships in a dataset is random.

Our major contributions include:

- 1. Making a relevant and usable connection between Ramsey theory of graphs and statistical analysis datasets (Definition 1).
- 2. Giving a statistical measurement of deviation from randomness (Definition 4).
- 3. Translating the Ramsey theorem Goodman's theorem to a measurement of transitivity of a system (Theorem 2).

In order for these connections to be further used and explored, we take care to explain the Ramsey theory we use in the language that an untrained data scientist will understand. Conversely, we take care to explain the statistics and data science in a way that is accessible to researchers in Ramsey theory.

In Section 2, we present the relevant definitions and mathematical framework. In Section 3, we introduce the needed Ramsey technology of Goodman's formula. In Section 4, we apply this to two real-life models: (1) similarity of voting records for the members of the 1984 US congress, and (2) economic trading data between countries. In Section 5, we give an application of Goodman's formula to measuring the transitivity of a graph. Finally, in Section 6, we discuss further directions for research.

2. Mathematical Framework

Here, we familiarize the reader with the notions related to graphs. If the reader is already familiar with these concepts, they may safely move to Section 3.

Our main model is a graph *G*, which is a collection of data points *V*, called the vertices, and a collection of connected (unordered) pairs of vertices *E*, called the edges, such that G = (V, E). An edge *a* between vertices v_1 and v_2 represents that v_1 and v_2 are related (in an abstract sense). This edge relationship will be intrinsic to each dataset and what it is trying to measure. For example, if the vertices are points in a metric space, we might assign an edge when the distance between two points is ≤ 1 . Another example is when the vertices are people in a room, and we put an edge between two people if they are mutual friends.

We insist that a vertex cannot be related to itself (a so-called loop) and that it can be described as an adjacency matrix by explicitly listing out which vertices have an edge between them:

A matrix $A = [a_{ij}]_{1 \le i,j \le N}$ is an adjacency matrix if it is symmetric with entries of 0, 1 with 0 s along the diagonal. An adjacency matrix can be thought of as a graph on vertices $\{1, ..., N\}$, where there is an edge between *i* and *j* iff $a_{ij} = 1$. This perspective is useful for the following reason:

Lemma 1. Let A be an $N \times N$ adjacency matrix, and $k \ge 1$. In the matrix $A^k = A \cdot A \cdot \ldots \cdot A$ (k-times) the *ij*th entry is the number of paths in A from *i* to *j* of length exactly k.

In other words, if the first power (k = 1) of the adjacency matrix A represents an edge (path length = 1) between two vertices v_1 and v_2 , higher powers of the adjacency matrix give us insight into the number of paths between v_1 and v_2 of length k. A graph with N vertices where all $\binom{N}{2}$ (pairwise) possible edges are included is called a complete graph, and is denoted by K_N . In the case N = 3, we call K_3 a triangle.

Corollary 1. Let A be an $N \times N$ adjacency matrix. The ii-th diagonal entry of A^3 is the number of triangles in A containing the vertex i. The number of triangles in A is $\frac{Trace(A^3)}{6}$, the sum of the diagonal entries of A^3 , taking into account over-counting.

Example of Corollary 1: Suppose we have a dataset with size N = 6. The number of triangles that exist in the complete K_6 graph is

$$\frac{\operatorname{Trace}(A(G_6)^3)}{6} = abf + acg + adh + aei + bcj$$
$$+ fgj + bdk + fhk + bel + fil$$
$$+ cdm + ghm + jkm + cen + gin$$
$$+ jln + deo + hio + klo + mno,$$

where each triplet (e_i, e_j, e_k) is a triplet of edges that create a triangle (K_3) . Depending on whether or not each edge has a value of 1 or 0 in the adjacency matrix *A* will determine if these triangles exist.

Suppose only the edges *a*, *b*, *c*, and *d* exist, as in Figure 1. Then, no triangles exist because when we replace the edges *a*, *b*, *c*, and *d* with 1 and everything else with 0, no triplet of edges is complete:

$$\frac{\operatorname{Trace}(A(G_6)^3)}{6} = (1)(1)(0) + (1)(1)(0) + (1)(1)(0) + (1)(0)(0) + (1)(1)(0) + (0)(0)(0) + (1)(1)(0) + (0)(0)(0) + (1)(1)(0) + (0)(0)(0) + (1)(1)(0) + (0)(0)(0) + (1)(0)(0) + (0)(0)(0)(0) + (0)(0)(0) + (0)(0)(0) + (0)(0)(0) + (0)(0)(0) + (0)(0)(0)$$

In this framework, if a triangle exists in the adjacency matrix A, then all three points (v_i, v_j, v_k) are connected to each other based on how the predetermined relationship is defined (whether it be geographic distance or some measurement of friendship, for example). In this way, a K_3 represents the simplest non-trivial emergent "pattern" that can be observed in a graph connecting data points in D, so it is the natural starting point for asking the question, "Which patterns are forced to exist in D given how we've connected its data points in the adjacency matrix A?".

This framework is good in black-and-white, binary situations where any pair of vertices is either (strongly) related or not related (at all). In non-binary relationships, it can be useful to think about graphs whose edges are classified by multiple colors. This can be represented as a partition of the edge set *E* into *r*-many disjoint sets $E = E_{c_1} \sqcup E_{c_2} \sqcup \ldots \sqcup E_{c_r}$, where c_1, \ldots, c_r represent a total of *r*-colors or classifications.



Figure 1. A graph where only the edges *a*, *b*, *c* and *d* exist. The dashed lines are used to indicate a lack of edge.

In the case of two colors, we will often just refer to red (R) and blue (B) edges. In the framework of adjacency matrices, a complete graph A with an edge-coloring using two colors is represented by an adjacency matrix B indicating a relationship exists or does not R:

$$B(G_6) = \begin{cases} v_1 & v_2 & v_3 & v_4 & v_5 & v_6 \\ v_2 & 0 & a & b & c & d & e \\ a & 0 & f & g & h & i \\ b & f & 0 & j & k & l \\ c & g & j & 0 & m & n \\ d & h & k & m & 0 & o \\ e & i & l & n & o & 0 \end{pmatrix},$$

$$B(G_6) = \begin{cases} v_1 & v_2 & v_3 & v_4 & v_5 & v_6 \\ v_1 & v_2 & v_3 & v_4 & v_5 & v_6 \\ 1-a & 0 & 1-f & 1-g & 1-h & 1-i \\ 1-b & 1-f & 0 & 1-j & 1-k & 1-l \\ 1-c & 1-g & 1-j & 0 & 1-m & 1-n \\ 1-d & 1-h & 1-k & 1-m & 0 & 1-o \\ 1-e & 1-i & 1-l & 1-n & 1-o & 0 \end{pmatrix}.$$

$$\therefore R(G_6) + B(G_6) = \begin{cases} v_1 \\ v_2 \\ v_3 \\ v_4 \\ v_5 \\ v_6 \end{cases} \begin{pmatrix} 0 & 1 & 1 & 1 & 1 & 1 \\ 1 & 0 & 1 & 1 & 1 & 1 \\ 1 & 1 & 0 & 1 & 1 & 1 \\ 1 & 1 & 1 & 0 & 1 & 1 \\ 1 & 1 & 1 & 1 & 0 & 1 \\ 1 & 1 & 1 & 1 & 0 & 1 \\ 1 & 1 & 1 & 1 & 1 & 0 \end{pmatrix}.$$

Take the edge *a* between v_1 and v_2 in *R* and set it equal to a = 1. Since the edge is colored red, it necessarily has to have an entry equal to zero (a - 1 = 1 - 1 = 0) in the blue edge adjacency graph B. In this case, R + B must be the matrix of all ones, except on the diagonal where it has zeros. Counting monochromatic triangles in *A* is particularly simple:

Corollary 2. Let A be an $N \times N$ adjacency matrix whose edges are colored using two colors. The number of monochromatic triangles in A is $\frac{\text{Trace}(B^3) + \text{Trace}(R^3)}{6}$.

Therefore, the total number of triangles in the dataset D is equal to the sum of red and blue triangles present in the adjacency matrices R and B. Notably, it only takes polynomial time to compute the number of single-colored (monochromatic) triangles in a graph.

3. The Ramsey Perspective

Classical Ramsey theory asks: "For given nonnegative integers *m*, *r*, does every edge coloring of a K_N complete graph with r colors contain a sub-collection K_m , all of whose edges have the same color?" In other words, how big does a multi-colored, complete graph need to be to force the existence of a smaller single-colored, complete graph?

In 1930, Ramsey [3] showed that if the size of the dataset D was $N \ge 6$, and the number of ways the data points could be related to each other was m = 2 (either related or unrelated), then unavoidable subgraphs of mutually related or unrelated data points are forced to exist.

In 1959, Goodman quantified how many single-colored (monochromatic) triangles must be present in a two-colored K_N . Because a K_3 represents the simplest object that describes how data points relate to each other beyond a simple edge, it will form the basis of our application of Ramsey theory.

Theorem 1 (Goodman 1959, [2]). Let G be a graph with N vertices and edge-colored with red and blue. The quantity of monochromatic triangles in G is at least:

- $\begin{array}{l} \frac{m(m-1)(m-2)}{3}, \ if \ N=2m,\\ \frac{2m(m-1)(4m+1)}{3}, \ if \ N=4m+1,\\ \frac{2m(m+1)(4m-1)}{3}, \ if \ N=4m+3. \end{array}$

Since the total number of triangles in K_N is $\binom{N}{3} = \frac{N(N-1)(N-2)}{6}$, Goodman's formula may be reinterpreted as a percentage.

Corollary 3 (Goodman 1959, [2]). Let G be a graph with N vertices and edge-colored with red and blue. The percentage of triangles in G that are monochromatic is asymptotically at least $\frac{N-3}{4N} \rightarrow \frac{1}{4}$.

This can be shown directly by dividing the quantities in Theorem 1 by $\binom{N}{3}$. Alternatively, by applying Schwenk's reformulation of Goodman's formula [4], we can easily prove this:

Proof. For *N* number of data points, the forced number F(N) of monochromatic red (*R*) and blue (*B*) triangles is:

$$F(N) = \binom{N}{3} - \left\lfloor \frac{1}{2}N \left\lfloor \frac{1}{4}(N-1)^2 \right\rfloor \right\rfloor,$$

and, since the number of triangles present in any complete graph is $\binom{N}{3}$, the following ratio describes the percentage of triangles in *G* that are monochromatic:

$$1 - \frac{\left\lfloor \frac{1}{2}N\left\lfloor \frac{1}{4}(N-1)^{2}\right\rfloor \right\rfloor}{\binom{N}{3}} = 1 - \frac{6\left\lfloor \frac{1}{2}N\left\lfloor \frac{1}{4}(N-1)^{2}\right\rfloor \right\rfloor}{(N-2)(N-1)N}.$$

The Floor Function of f(x) is equivalent to the function of f(x) with discontinuities at non-integer values x, therefore describing the asymptotic nature of the above ratio can be done without taking the floor functions into consideration:

$$1 - \frac{6\frac{1}{2}N(\frac{1}{4}(N-1)^2)}{(N-2)(N-1)N} = \frac{1}{4} - \frac{3}{4(N-2)} \therefore \lim_{n \to \infty} \frac{1}{4} - \frac{3}{4(N-2)} = \frac{1}{4}.$$

From this, we can establish a threshold for when a two-colored graph can be interpreted to have meaningful correlations.

Definition 1. Let G be a graph with n vertices and edge-colored with red and blue. Let Mono(G) be the percentage of triangles in G that are monochromatic, among all possible $\binom{N}{3}$ triangles in G. Let Goodman(N) be the minimum percentage of monochromatic triangles in G guaranteed by Corollary 3, which has been shown to approach 0.25 as $N \to \infty$. If Mono(G) > Goodman(N), then we say that G has potentially meaningful correlations, which we explore further in Section 4.2.

If Mono(G) is much larger than Goodman(N), then we might say that *G* obeys a triangle dichotomy, which means that we expect a lot of triangles to be either completely one color, or completely the other. This is a relative term, and the larger Mono(G) is the more that this resembles a true dichotomy. If one color is more heavily represented in *G* than another, then we might say that *G* has a triangle bias. When triangle bias exists, this is at odds with the expectation, in a randomly colored graph, that the ratio of the number of color R triangles to color B triangles should be 1:1, and is therefore a further indication that the correlations in *G* are meaningful.

How can this be used in a dataset? In Section 4.2, we discuss a best-fit approach in order to test the null-hypothesis that a dataset is indeed random. In the next section, we show how diverse datasets can be modeled by graphs.

4. Models

Many datasets come naturally as tuples of variables of data, not as a graph. Here, we give a meaningful way to capture information about similarity in the data using a graph. At the most basic level, any measurement of how similar two variables are translates into a measurement of how similar two tuples of variables are (such as by taking an average, or a weighted average).

We will focus our analysis on two datasets: (1) voting records of members of the US congress in 1984, and (2) economic partnership among countries. In the first dataset, we are able to vary our way of combining the information about how the individual variables are similar using what we call threshold graphs. These variations give us a way to very closely compare how far the dataset deviates from truly random. In the second dataset, we do not have complete information, but we are still able to give a more limited measurement of this deviation.

4.1. Similarity in Voting Records

We now look at a set of people that have voted multiple times, specifically the 1984 United States Congressional Voting Records [5]. There is no special motivation for picking this specific dataset, or year, other than it was available to the public. We translate this dataset into a graph with 435 nodes and $\binom{435}{2}$ labeled edges.

Goodman's formula can quantify how strong the triangle dichotomy and triangle bias are; that is, the percentage of three person cliques (B) and independent triples (R) and their deviation from the expected 1:1 color ratio. We will use the Hamming distance to measure how similar two voting records are for the 435 congress members of the 1984 US congress.

Definition 2. *The Hamming distance of two strings of the same length is the total number of positions where the entries are different.*

For example, the Hamming distance between 00010 and 01001 is 3. These strings differ in the second, fourth and fifth spots.

In this session, there were 16 separate votes, and to each voter we assign the string of length 16 with entries 'N' (voted nay), 'Y' (voted yea) or 'A' (some other action, such as abstaining). The minimum Hamming distance is 0, which indicates two identical voting records, and the maximum distance is 16, meaning the two voters always voted differently. See Table 1 for sample data.

Voter	Party		Voting String		
v_1	R	NYNY	YYNN	NYAY	YYNY
v_2	R	NYNY	YYNN	NNNY	YYNA
v_3	D	AYYA	YYNN	NNYN	YYNN
v_4	D	NYYN	AYNN	NNYA	YNNY
v_5	D	YYYN	YYNN	NNYA	YYYY
v_6	D	NYYN	YYNN	NNNN	YYYY

Table 1. The voting entries of the first six voters in the 1984 US congress dataset.

Applying this notion of distance to the data from Table 1 gives the following adjacency matrix and graph (Figure 2):

	v_1	v_2	v_3	v_4	v_5	v_6
v_1	(0	3	7	7	7	6 \
v_2	3	0	6	7	7	5
v_3	7	6	0	5	5	5
v_4	7	7	5	0	5	4
v_5	7	7	5	5	0	3
v_6	6	5	5	4	3	0/



Figure 2. The graph that corresponds to the first six voting records in the 1984 US Congress dataset.

How do we turn this into a binary adjacency matrix for classification purposes? In other words, how do we decide what constitutes "similar" voting and "dissimilar" voting, and where do we make that cut off?

Definition 3. Let (M, d) be a metric space with vertex set M and distance d. Let $t \ge 0$. Define the two-colored threshold graph G(t) in the space (M, d) by coloring an edge between two vertices v_i, v_j blue B iff $d(v_i, v_j) > t$, and red R iff $d(v_i, v_j) \le t$.

Therefore, when for example t = 5, the following graph G(t = 5) maps to the binary case:

	v_1	v_2	v_3	v_4	v_5	v_6			v_1	v_2	v_3	v_4	v_5	v_6
v_1	(0	3	7	7	7	6 \	١	v_1	(0	0	1	1	1	1 \
v_2	3	0	6	7	7	5		v_2	0	0	1	1	1	0
v_3	7	6	0	5	5	5	P(C) =	v_3	1	1	0	0	0	0
v_4	7	7	5	0	5	4	$\rightarrow D(G_6) =$	v_4	1	1	0	0	0	0
v_5	7	7	5	5	0	3		v_5	1	1	0	0	0	0
v_6	6	5	5	4	3	0 /	/	v_6	$\backslash 1$	0	0	0	0	0/
	v_1	v_2	7/2	71	~ 1				714	720	710	~1	71-	71
		-	03	v_4	v_5	v_6			v_1	υZ	03	v_4	v_5	06
v_1	(0	3	7	$\frac{\partial_4}{7}$	05 7	$\begin{bmatrix} v_6 \\ 6 \end{bmatrix}$		v_1	$\begin{pmatrix} 0 \\ \end{pmatrix}$	1	0	0^{0}_{4}	05 0	0
v_1 v_2	$\begin{pmatrix} 0\\ 3 \end{pmatrix}$	3 0	7 6	04 7 7	05 7 7	$\begin{pmatrix} v_6 \\ 6 \\ 5 \end{pmatrix}$		$v_1 \\ v_2$	$\begin{pmatrix} 0\\ 1 \end{pmatrix}$	1 0	0 0 0	0 0	0 0 0	$\begin{pmatrix} 0 \\ 1 \end{pmatrix}$
v_1 v_2 v_3	$\begin{pmatrix} 0\\ 3\\ 7 \end{pmatrix}$	3 0 6	7 6 0	7 7 7 5	05 7 7 5	$\begin{bmatrix} v_6 \\ 6 \\ 5 \\ 5 \end{bmatrix}$	$\mathcal{P}(C_{i}) =$	v_1 v_2 v_3	$\begin{pmatrix} 0\\ 1\\ 0 \end{pmatrix}$	1 0 0	0 0 0		0 0 1	$\begin{pmatrix} 0 \\ 1 \\ 1 \end{pmatrix}$
v_1 v_2 v_3 v_4	(0 3 7 7	3 0 6 7	7 6 0 5	7 7 5 0	05 7 7 5 5	$v_6 \\ 6 \\ 5 \\ 5 \\ 4$	$\rightarrow R(G_6) =$	v_1 v_2 v_3 v_4	$\begin{pmatrix} 0\\ 1\\ 0\\ 0 \end{pmatrix}$	1 0 0 0	0 0 0 1	$ \begin{array}{c} 0_4 \\ 0 \\ 0 \\ $	0 0 1 1	$\begin{pmatrix} 0 \\ 1 \\ 1 \\ 1 \\ 1 \end{pmatrix}$
$ \begin{array}{c} v_1 \\ v_2 \\ v_3 \\ v_4 \\ v_5 \end{array} $	(0 3 7 7 7 7	3 0 6 7 7	7 6 0 5 5	7 7 5 0 5	7 7 5 5 0	v_6 5 5 4 3	$\rightarrow R(G_6) =$	$v_1 \\ v_2 \\ v_3 \\ v_4 \\ v_5$	$ \begin{pmatrix} 0 \\ 1 \\ 0 \\ 0 \\ 0 \end{pmatrix} $	1 0 0 0 0	0 0 0 0 1 1	$ \begin{array}{c} 0_4 \\ 0 \\ 1 \\ $	0 0 1 1 0	$\begin{pmatrix} 0 \\ 1 \\ 1 \\ 1 \\ 1 \\ 1 \end{pmatrix}$

For example, taking the sample Congressional voting data from Table 1, we get the following threshold graphs in Figures 3 and 4.



Figure 3. The graphs G(t = 0), G(1), G(2) (left), G(3) (center), and G(4) (right).



Figure 4. The graphs G(5), G(6) and G(7), ..., G(17).

Consider the complete graph G(t), the subgraph composed entirely of Democrat Congress votes D(t), and the subgraph composed entirely of Republican Congress votes R(t). Applied to the total voting records available in the dataset at various thresholds t, the following table (Table 2) and figure (Figure 5) show the ratio of Mono(G(t)) to the total number of triangles K_N :

t	G(t)	D(t)	R(t)	t	G(t)	D(t)	R(t)
0	1.000	1.000	1.000	9	0.271	0.496	0.770
1	0.993	0.933	0.972	10	0.299	0.586	0.842
2	0.953	0.953	0.829	11	0.370	0.688	0.891
3	0.858	0.850	0.603	12	0.471	0.783	0.932
4	0.727	0.699	0.475	13	0.597	0.871	0.954
5	0.590	0.549	0.461	14	0.743	0.943	0.964
6	0.462	0.440	0.506	15	0.888	0.979	0.977
7	0.359	0.399	0.581	16	0.970	0.997	0.984
8	0.291	0.423	0.672	17	1.000	1.000	1.000
Goodman	0.248	0.247	0.246	Goodman	0.248	0.247	0.246

Table 2. The percentage of monochromatic triangles for various threshold graphs. The minimum values are boxed.



Figure 5. Data from Table 2 visualized.

In Section 5, we give another natural interpretation of these results by giving a measure of how transitive these graphs are. This is maybe a more intuitive interpretation of the data since it gives us a direct measurement of cooperation and independence.

4.2. How to Measure the Deviation Away from a Random Graph

Goodman's formula tells us how many monochromatic triangles are forced to exist for a dataset D of size N, but what would the threshold graph of a truly random coloring of G_N look like?

4.2.1. Theoretical Construction

Supposing that we have a random graph G(N, t) of related to N data points in D and probability t that an edge between two vertices n_i, n_j exists, the Erdos–Renyi model tells us that the expected number of edges in G(N, t) is $\binom{N}{2}t$. The parameter t can be thought of as the threshold parameter introduced in Section 4.1 as it ranges from $0 \rightarrow 1$ (assuming t_{min} and t_{max} have been normalized to [0, 1]). Therefore, the expected number of red (R) and blue (B) triangles T in D is:

$$\mathbb{E}[T^R] = \binom{N}{3}(t^3), \mathbb{E}[T^B] = \binom{N}{3}(1-t)^3,$$

where $\binom{N}{3}$ represents the total possible number of triangles, and t^3 represents the probability that all three edges are red (*R*); likewise, $(1 - t)^3$ represents the probability of all three edges being blue (*B*). This information can be used to calculate the number of monochromatic triangles.

Corollary 4. The expected number of monochromatic triangles in G_N is $2\binom{N}{3}(\frac{1}{2})^3$.

Proof. The probability that three adjacent edges are the same color in a 2-colored graph is $(\frac{1}{2})^3$, there are $\binom{N}{3}$ number of triangles, and we multiply by 2 to account for the symmetry of how the edges can be colored with equal probability. \Box

This creates the following threshold plot (Figure 6) for any randomly colored graph G(N, t).



4.2.2. Defining Deviation

Definition 4. Deviation is a combination of the degree of (A): the triangle dichotomy and (B): the triangle bias.

Deviation away from the expected distribution can allow us to determine the likelihood that the null hypothesis H_0 (that G_N is actually random) is accepted or rejected. This can be done with a simple χ^2 test:

$$\chi_R^2 = \sum_{i=t_{min}}^{t_{max}} \frac{(R(i)_O - E[R]_i)^2}{E[R]_i}, \chi_B^2 = \sum_{i=t_{min}}^{t_{max}} \frac{(B(i)_O - E[B]_i)^2}{E[B]_i}.$$

The average of χ_R^2 , χ_B^2 and their resulting *p*-value can be used to determine with some significance level whether to accept or reject H_0 .

While the expected value is a good benchmark, it still doesn't answer the more fundamental question of how many monochromatic triangles are present in G_N versus how many are required by

Ramsey theory. This creates a stricter χ^2 calculation, but one that's better suited to our needs and is a measurement of the triangle dichotomy and triangle bias:

$$\chi_G^2 = \sum_{i=t_{min}}^{t_{max}} \frac{(G(i)_O - F(N))^2}{F(N)},$$
$$\chi_R^2 = \sum_{i=t_{min}}^{t_{max}} \frac{(R(i)_O - \frac{F(N)}{2})^2}{\frac{F(N)}{2}}, \\ \chi_B^2 = \sum_{i=t_{min}}^{t_{max}} \frac{(B(i)_O - \frac{F(N)}{2})^2}{\frac{F(N)}{2}}.$$

4.2.3. Applied to Voting Threshold Graphs

We are now faced with applying the χ^2 method from Section 4.2.2 to the Congressional voting threshold graphs. What is the likelihood that these are random, or, equivalently, what is the likelihood that there is a bias in the congressional voting record? This is answered in Tables 3 and 4.

Table 3. The χ^2 fit for the overall voting record G(t), Democrats D(t), and Republicans R(t). This demonstrates the degree of the triangle dichotomy for each pre-defined classification.

Quantity	G(t)	D(t)	R(t)
Goodman	0.248	0.247	0.246
χ^2	17.448	22.552	25.206

Table 4. The χ^2 fit for the overall voting record G(t), Democrats D(t), and Republicans R(t) by color (R,B). This demonstrates the degree of the triangle bias for each pre-defined classification.

Subset	Blue χ^2	Red χ^2	Total χ^2
Total $G(t)$	18.7782	22.864	17.448
Democrats $D(t)$	28.536	22.596	22.552
Republicans <i>R</i> (<i>t</i>)	22.705	38.028	25.206

These χ^2 values have *p*-values that are very, very small. A way to place these in context is to compare them to the expected value's deviation from what's required by Ramsey theory. See Tables 5–7.

Table 5. The χ^2 fit for the overall expected value of forced monochromatic triangles.

Quantity	Blue χ^2	Red χ^2	Total χ^2
Expectation	16.384	16.384	10.076

Table 6. The deviation of χ^2 of G(t), D(t), and R(t) from their respective expected χ^2 values.

Subset	Blue χ^2	Red χ^2	Total χ^2
Total	18.7782 - 16.384 = 2.394	6.48	17.448 - 10.076 = 7.372
Democrats	12.152	6.212	12.476
Republicans	6.321	21.644	15.130

Table 7. The *p*-value associated with each of these is based on the cumulative distribution function (CDF), namely p = 1 - CDF. At a significance level of 0.01, the non-significant deviations are <u>underlined</u>.

Subset	Blue χ^2	Red χ^2	Total χ^2
Total	0.121802	<u>0.010909</u>	0.006625
Democrats	0.00049	0.012689	0.000412
Republicans	0.011932	< 0.00001	0.0001

We can say a *p*-value is significant if it is sufficiently different from how the expectation value differs from what is required from Ramsey theory. At a significance level of 0.01, the non-significant deviations in Table 7 are <u>underlined</u>. (The significance level selected depends on the cost function of the particular model. For instance, the significance level would scale with the cost associated with being wrong.). The furthest deviation can be attributed to the Republican congressional voters and is an indication that a strong bias exists in their voting records.

4.3. Collaboration Model

Suppose we have a collection of people V, working together on a communal project.

As an example, we look at economic trading data [6,7]. Every country is represented by a node, and we add a blue edge from a country to its five largest importers and exporters by volume. See Figure 7 for a visulaization. In this way, two countries are connected by a blue edge if their countries are historically economically connected and by a red edge if they are smaller trading partners. There is an asymmetry in the way edges are added, as, for example, China only adds at most 10 blue edges to other countries, but many countries add blue edges to China. In this way, it is possible for a country to have a blue degree much higher than 10. This graph is best described as an Interaction Graph similar to the "friends at a party".



Figure 7. Countries are arranged alphabetically starting at the top and going counterclockwise. The green nodes are the G7 and G20 countries. The graph has 214 vertices, 1363 blue edges, the average blue degree is 12.7, the five highest blue degrees are 162 (China), 125 (United States), 96 (Germany), 66 (France) and Italy (61). The largest complete subgraph has eight vertices: Algeria, China, France, Germany, Italy, Spain, United Kingdom, and the United States, forming a K_8 . The largest independent set has 70 vertices, forming an I_{70} .

For N = 214 countries, the number of monochromatic triangles equals 85.0% of the total number $\binom{N}{3}$ of triangles in a K_{214} . These monochromatic triangles are almost entirely red, representing a lack of strong trade relations. This is significantly more than the required number of triangles given by Goodman's formula, which at N = 214 is 24.7%. See Table 8.

Table 8. Percentages of monochromatic triangles for the country graph.

Quantity	Percentage
Percentage of red K_3	84.8%
Percentage of blue K_3	0.2%
Percentage of mono-chromatic triangles	85.0%
Goodman-type lower bound	24.7%

Since this graph has a threshold of only the top five trading partners for each country, it can be seen as a discrete sample of the threshold graph that would exist on the scale (t_{min} = top trading partner to t_{max} = all trading partners). In order to determine if the percentage of monochromatic triangles in this graph can be interpreted as meaningful evidence that the global economy connected with a strong dichotomy, we need to measure its *p*-value. For *n* = 214 countries, a threshold of *t* = 5 corresponds to *t* = 0.0234 on a normalized scale of [0, 1]. When *t* = 0.0234, the expected deviation for the total number of monochromatic triangles from those required by Ramsey theory has a χ^2 = 4.236, whereas the trading graph has a χ^2 = 2.907. The difference between these is 1.329, which corresponds to a *p*-value of 0.248983, which is not statistically significant. We can therefore not reject the null-hypothesis that this trade graph is random.

While we cannot reject H_0 based on the number of superfluous monochromatic K_3 's in the trading data, the presence of higher dimensional complete subgraphs might provide sufficient evidence.

We can compute the percentage of monochromatic K_4 , and the percentage of monochromatic K_5 . This is computationally complex, so we computed these percentages (Table 9) for only small N.

N	Percentage of Mono-Chromatic K_N (%)	Goodman-Type Lower Bound
3	85	25
4	74	3
5	62	<1

Table 9. Data for the country graph in Section 4.3.

It is natural to then ask what happens when we consider larger substructures, that is, $K_4, K_5, ..., K_N$ instead of triangles.

χ^2 for Higher Dimensions

For higher dimensions, there is no analogue of Goodman's formula, which we would expect to give us a percentage of $\frac{1}{32}$ for K_4 , $\frac{1}{16384}$ for K_5 , etc. using the same methods described in Corollary 4. See Figure 8. In [8], Thomason has shown that an upper bound for the corresponding percentage of monochromatic K_4 is $\frac{1}{33}$, although it is not known if this is tight. In the same work, he gave an upper bound on the number of monochromatic K_m , as $0.936 \cdot 2^{1-\binom{m}{2}}$.



Figure 8. The expected number of monochromatic K_4 and K_5 s as a function of t. The Goodman-type upper bound for K_4 is 0.0295, and 0.00183 for K_5 .

For the χ^2 's related to larger substructures, Thomason's upper bound can be used in the same way that Goodman's is used for K_3 , with the understanding that this will give us an upper bound on a graph's deviation from what is required by Ramsey theory. Our new $\bar{\chi}^2$ is an average of each K_m 's associated χ^2 and can include up to *N*-dimensional substructures:

$$\bar{\chi}^2 = rac{1}{N-3} \sum_{i=3}^N \chi^2_{K_i}.$$

If instead we increase the number of colors and therefore allow for more than two classifications, a perfect answer for three colors and triangles is given by [9].

4.4. Applications to Other Datasets

We end this section with a more general discussion about using these methods on different datasets.

The relevant part of the voting dataset was that each data point (voter) was a list of variables (votes), and we had a measurement of what it means for two entries of the same variable to be similar. In the case of voting records, our measurement of similarity was very blunt: either they were the same or they were different.

This method of analysis could be applied to datasets where the variables have different measures of similarities, such as comparing the background of two potential clients for an insurance company. In this case, for example, our measurement for similarity in age might be simply a difference of the two ages, but the measurement of similarity of education levels would be more subtle.

The methods presented here give a measurement of how far a dataset deviates from random. If, for example, a company wishes to build a diverse, nearly random roster of clients, this can be used to quantify how close to that goal the company is.

5. Applications to Transitivity

When we have sufficient evidence to reject H_0 , we define a non-random graph in terms of its transitivity. Transitivity can be thought of as the likelihood that a relationship in a dataset is meaningful and therefore not spurious. Let's again consider the model for the party problem: the nodes are people at a party and we assign a blue (*B*) edge between two people if they are friends (and red (*R*) if they are not friends).

Definition 5. A binary relation R on D is transitive if $\forall v_i, v_j, v_k \in D$ if $v_i R v_j$ and $v_j R v_k$ then $v_i R v_k$.

In this setting, we first remark that the blue "friend" relation is not by-default transitive, and neither is the red "not friend" relation. For example, I am friends with someone who does not know my brother.

It is easy to see that the only way for the red relation to be transitive is if *all* edges are red in a particular subgraph. Similarly, the blue relation is transitive only if all edges are blue. Typically, such a graph will not be transitive in both relations.

Transitivity can be described in terms of monochromatic triangles, specifically three vertices v_i, v_j, v_k are members of a graph that is not transitive when the edges between them are not monochromatic. In this way, the percentage of monochromatic triangles in a graph is a measure of how transitive a graph is. In the context of uncolored graphs, this has been studied as the clustering coefficient. However, by looking at two colored graphs, Goodman's formula implies that there is a lower limit on how non-transitive a graph can be. We know that least 0.25 of its triangles must be monochromatic in the case of a two colored graph. The higher the observed percentage is than 0.25, the more transitive the graph is, and this can be measured in terms of χ^2 .

Let's use this to interpret the results from Section 4.1. Suppose we have three democrats v_i , v_j , v_k and we know that $v_i R v_j$ iff $v_j R v_k$; that is, the relationship between v_i and v_j is the exact same as the one between v_i and v_k (although we don't necessarily know if both have an edge or not).

We ask: how likely is it that the relationship between v_i and v_j is the same as the one between v_i and v_k , i.e., that the triangle is transitive?

Theorem 2. Let G be a complete graph with N vertices whose edges are colored red (R) or blue (B). The percentage of monochromatic paths of length 2 that complete to a monochromatic triangle is measured by

$$\frac{3f(G)}{\binom{N}{3}+2f(G)},$$

where f(G) is the number of monochromatic triangles in G.

Proof. This quantity comes from the observation that every monochromatic triangle contains three monochromatic paths of length 2, but each non-monochromatic triangle contains precisely one monochromatic path of length 2. For ease of computation, we use that (the number of non-monochromatic triangles) + $3 \times$ (the number of monochromatic triangles) is $(\binom{N}{3} - f(G)) + (3f(G)) = \binom{N}{3} + 2f(G)$, since $\binom{N}{3}$ is the total number of triangles. Thus, $\binom{N}{3} + 2f(G)$ is the total number of monochromatic triangle paths of length 2 in *G*, since this counts every non-monochromatic triangle once and counts every monochromatic triangle three times. \Box

By using Goodman's formula, this observation above translates to the following (completely expected) result:

Proposition 1. Let *G* be a graph with *N* vertices and edge-colored with red and blue. The ratio of monochromatic paths in G that are part of a monochromatic triangle is asymptotically at least 0.5.

The observation above provides an efficient way to compute the ratio of monochromatic paths in *G* that are part of a monochromatic triangle. We, for example, don't need to count the number of monochromatic paths directly.

5.1. Application to Previous Examples

5.1.1. Application to Voting Records

In the case of the threshold graphs from Section 4.1, the threshold graph G(t) with the minimum "transitivity percentage" is precisely the threshold graph with the minimum number of monochromatic triangles, namely t = 9 (52.7%). Analogously, for D(t), this occurs at t = 7 (66.6%) and for R(t) this occurs at t = 5 (72.0%). See Table 10 and Figure 9.

Table 10. 7	Fransitivity num	bers for the th	reshold graphs.	The minimum	values are b	oxed.

t	G(t)	D(t)	R(t)	t	G(t)	D(t)	R(t)
0	1.000	1.000	1.000	9	0.526	0.747	0.909
1	0.997	0.997	0.990	10	0.561	0.809	0.941
2	0.984	0.983	0.935	11	0.637	0.868	0.960
3	0.947	0.944	0.819	12	0.727	0.915	0.976
4	0.888	0.874	0.730	13	0.816	0.952	0.984
5	0.811	0.785	0.719	14	0.896	0.980	0.987
6	0.719	0.701	0.754	15	0.959	0.993	0.992
7	0.626	0.665	0.806	16	0.989	0.998	0.994
8	0.552	0.687	0.859	17	1.000	1.000	1.000



Figure 9. Transitivity percentages for the graphs G(t), D(t) and R(t), with $0 \le t \le 17$.

5.1.2. Application to Global Trading Data

China provides an interesting example of a country that is part of many non-monochromatic triangles because China has an edge to 162 of the 213 other countries, and of those 162 countries only 6.9% of edges are present among China's neighbors. Thus, only a small percentage of China's neighbors are themselves directly connected. This contributes to slightly lowering the percentage of transitivity in the larger graph.

In total, using all countries, 94.4% of all monochromatic paths complete to an edge of the same color. This is well above the 50% guaranteed by Proposition 1. Again, a complication is introduced by only looking at one threshold level rather than calculating the entire χ^2 .

6. Conclusions and Questions

We now make two major calls to use these methods: applications and development of related theory.

6.1. Theory Building

This use of Goodman's formula suggests the need for other quantitative Ramsey statements. For higher dimensional objects, we mention a couple that already exist and some that have yet to be developed.

A recent survey of Ramsey bounds for hypergraphs is a useful place to see the current best known bounds for various Ramsey numbers [10]. This survey also goes through proof sketches, many of which contain a weak Goodman-style lower bound. These bounds typically come from a use of the probabilistic method (see, for example, [11]).

In general, the probabilistic bounds provide a first non-trivial upper bound on the percentage of monochromatic structures, and improving them can be difficult. In order to use Ramsey theory in a generalized way, a closed form analogous to Goodman's formula needs to be developed for all K_n subgraphs and all C_n -colored graphs.

6.2. Further Applications

The case of triangles is simple, but still captures the quantitative notion of transitivity of a relation. Additionally, counting the number of monochromatic triangles in a graph is computationally efficient.

Further progress could be motivated by finding interpretations for other quantitative Ramsey statements. For example, a quantitative version of Van der Waerden's theorem for a fixed length. That is, given a 2-coloring of the points $\{1, 2, ..., 9\}$, it is known that there must be at least one arithmetic progression of length 3 (i.e., $a_0, a_0 + m, a_0 + 2m$) where all points are the same color. The following question has a reasonable answer in [12], which has serious mathematical content:

Remark 2 (Question). For N sufficiently large. Give reasonable lower-bounds and upper bounds on the percentage of monochromatic 3-term progressions that must exist for any 2-coloring of $\{0, 1, 2, ..., n\}$.

In [12], it is shown that, asymptotically, at least 25% of all 3-term such arithmetic progressions must be monochromatic. This extended results of [13]. In their setting, arithmetic progressions are allowed to "wrap around". That is, in $\{0, 1, 2, 3, 4, 5, 6, 7\}$, the triple $\{5, 7, 1\}$ is considered a 3-term progression.

For 4-term progressions, see [14] and the strengthening [15]. Both of these are non-trivial results.

The next step is to interpret 3-term progressions (or 4-term progressions) in a data-set in a meaningful, physical way.

6.3. Closing Remarks

We believe that the connections between data science and Ramsey theory are still largely unmade and will prove to be profound. We have shown that Ramsey theory can be used to rigorously define spurious correlations in datasets, and how deviations from the number of required spurious correlations might be meaningful in terms of transitivity.

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