Article

# Statistical Analysis of Maximally Similar Sets in Ecological Research

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Abstract: Maximally similar sets (MSS) are sets of elements that share a neighborhood in a high-dimensional space defined by symmetric, reflexive similarity relation. Each element of the universe is employed as the kernel of a neighborhood of a given size (number of members), and elements are added to the neighborhood in order of similarity to the current members of the set until the desired neighborhood size is achieved. The set of neighborhoods is then reduced to the set of unique maximally similar sets by eliminating all sets that are permutations of an existing set. Subsequently, the within-MSS variability of attributes associated with the elements is compared to random sets of the same size to estimate the probability of obtaining variability as low as observed. Individual attributes can be compared for effect size by the ratio of within-MSS variability to random set variability, correcting for statistical power as necessary. The analyses performed identify constraints, as opposed to determinants, in the triangular distribution of pair-wise element similarity. In the example given here, the variability in spring temperature, summer temperature, and growing degree days of forest vegetation samples shows the greatest constraint on forest composition of a large set of candidate environmental variables.

**Keywords:** similarity relation neighborhoods, similarity relation decomposition, statistical analysis of within-set variability

#### 7 1. Introduction

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The discipline of community ecology investigates variability in the composition of ecological communities and the possible environmental factors that might determine or constrain that composition. Because ecological communities generally contain multiple species the comparison of composition is inherently multivariate. The general approach to the analysis of this variability involves the calculation of a symmetric matrix of similarities, dissimilarities, or distances between all possible pairs of community samples, followed by the subsequent analysis of the properties of that matrix. This analysis typically takes one of two forms: partitioning the sample units into sets of similar samples through some form of cluster analysis, or projecting the matrix to lower dimensionality and analyzing the variability as a field.

Formal statistical analysis of ecological communities goes back to at least 1954, when David Goodall introduced the use of Principal Components Analysis (PCA) to ecology [1]. Over the subsequent decades numerous statistical methods have been adopted or invented to further the aims of community ecology. In particular, various forms of cluster analysis have been investigated and compared for ecological analysis. Recent examples of comparative analyses of clustering algorithms applied to ecological community data are given by Roberts [2] and Aho et al. [3]. Generally, if not without exception, ecological cluster analyses have produced partitions of the sample units, i.e. a family of sets where (1) every set has at least one member, (2) every element belongs to exactly one set, and (3) the union of the sets comprises the universe of elements. While there are practical reasons to desire a partition as the outcome of the analysis (e.g. to produce a set of community types for inventory

or mapping purposes) the use of partitions presents challenges in statistical analysis of the underlying environmental constraints or determinants of the community type composition.

Given a sample with *N* sample units, the number of possible partitions of the data into clusters is given by Bell's number

$$B_n = \sum_{k=0}^{N} \begin{Bmatrix} N \\ k \end{Bmatrix}. \tag{1}$$

where k is a given number of clusters ( $0 \le k \le N$ ), N is the number of sample units, and  $\begin{Bmatrix} N \\ k \end{Bmatrix}$  is the Stirling number of the second kind.

$${\binom{N}{k}} = \frac{1}{k!} \sum_{j=0}^{k} (-1)^{k-j} {\binom{k}{j}} j^{N}.$$
 (2)

While the population of possible partitions is generally large, sampling at random from that population is difficult due to the peculiarities of the cluster analysis result to be compared to. Even constraining the sample population to the same value of *k* it may or may not be desirable to constrain the individual clusters to the distribution of cluster sizes obtained in the original cluster analysis, which is in part an artifact of the particular cluster analysis performed and not necessarily a good result.

Alternatively, it is possible to calculate a covering, as opposed to a partition of the data, i.e. a family of sets such that (1) all sets have at least one member, (2) all elements belong to at least one set, and (3) the union of all sets comprises the universe of sample units. By relaxing the partition constraint that every element belongs to exactly one set, but imposing the constraint that all sets are the same size, we can derive a covering, as opposed to a partition. In this case the number of possible sets of size  $n_k$  in the covering for k clusters is

$$\binom{N}{n_k}$$
 (3)

where  $n_k$  is the number of elements of set k. This is a much more favorable population to sample from. We simply sample  $n_k$  sample units without replacement from the set of N sample units a large number of times and compare the random samples to the the observed sets in the covering.

# 57 2. Materials and Methods

## 58 2.1. Data

The proposed analysis is demonstrated on a sample of the forest vegetation of the Shoshone
National Forest, Wyoming, USA. Sample units were 375 m<sup>2</sup> circular plots where the abundance of
every vascular plant species was estimated according to an ordinal ten-class scale. Environmental
attributes associated with the sample units were either measured in the field or modeled in a geographic
information system. Attributes include sample unit elevation, aspect, slope steepness, surficial geology,
soil properties, topographic position, and climate attributes modeled from elevation, aspect, slope,
surficial geology, topographic position, and geographic location. One hundred fifty sample units were
selected at random from a larger study to provide a manageable example data set. Further details
about the data are given in Appendix A.

## 68 2.2. Analyses

The sample unit composition data were used to calculate a symmetric, reflexive similarity matrix using the Bray-Curtis index [4].

$$s_{ij} = \frac{\sum_{q=1}^{m} 2 \times \min(a'_{iq}, a'_{jq})}{\sum_{q=1}^{m} a'_{iq} + a'_{jq}}$$
(4)

where  $s_{ij}$  is the similarity of sample unit i to sample unit j, m is the number of species,  $a_{iq}$  is the abundance of species q in sample unit i, and  $a'_{iq} = log(a_{iq} + 1)$ .  $s_{ij} \rightarrow [0, 1]$  where sample units with no species in common = 0 and identical sample units = 1.

Maximally similar sets (MSS) were solved for by setting the desired neighborhood size  $n_k$ , and then iteratively for each sample unit adding the sample unit most similar to the members of the neighborhood until the desired neighborhood size was achieved. The most similar sample unit to the neighborhood was calculated as

$$s_{ik_x} = \max \min_{\substack{i=1\\i\ni k_x; j\in k_x}}^N s_{ij} \tag{5}$$

where  $s_{ik_x}$  is the similarity of sample unit i to neighborhood  $k_x$ .

The set of resulting neighborhoods was reduced to the set of unique neighborhoods by deleting all neighborhoods that were a permutation of an existing neighborhood. The number of neighborhoods in the similarity relation (*S*) reflects the topology of the similarity relation and cannot be known *a priori*. As the size of neighborhoods goes up the number of neighborhoods generally declines.

The within-MSS variability of interval- or ratio-scaled sample unit attributes was examined by calculating the range of the attribute within each MSS. That set of *K* ranges was then compared to the set of ranges of an equal number of sets of equal size drawn at random without replacement from the set of sample units. The ranges of the MSS were compared to the ranges of the randomly drawn sets in a Wilcoxon rank-sum test with continuity correction to generate the Wilcoxon statistic *W*. The within-MSS variability for categorical variables was examined by calculating the entropy of the tabulated values of the variable for each the *K* MSS compared to the entropy of *K* sets of the same size sampled at random without replacement from the set of sample units. The entropy for set *k* was calculated as

$$e_k = -\sum_{\substack{c=1\\n_c \neq 0}}^{C} p_c \times \log(p_c); \quad p_c = n_c/N$$
(6)

where  $n_c$  is the number of sample units in category c for categorical variable C. As for the interval-valued attributes, the observed and random entropies were then tested with a Wilcoxon rank sum test with continuity correction to generate the Wilcoxon statistic W. This process was repeated 1000 times for each attribute to generate 1000 W statistics for each attribute, and the effect size of each attribute was estimated by comparing the distribution of W values in a boxplot.

### 3. Results

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# 3.1. Similarity Relation and Maximally Similar Sets

The distribution of similarities in the Bray-Curtis similarity relation is shown in Figure 1. Figure 1a shows relatively few cases of  $s_{ij} = 0$ , and relatively few cases of  $s_{ij} > 0.4$ . Figure 1b shows that mean similarity and minimum similarity are somewhat correlated, and that most sample units have at least

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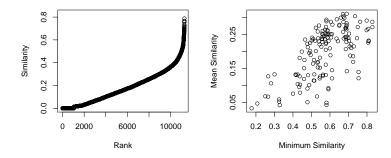
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**Figure 1.** Distribution of similarity values in the Bray-Curtis similarity relation. (a) The cumulative empirical density distribution. (b) The pairwise distribution of minimum and mean similarity for each sample unit.

one other sample unit with  $s_{ij} > 0.5$ . This is a typical result for vegetation data where the composition of the sample units is quite diverse but where sampling is adequate to span the range of variability.

MSS were calculated for neighborhood sizes of 5, 10, 15, and 20 members. Table 1 gives the sizes of the resulting families of neighborhoods.

**Table 1.** Number of neighborhoods as a function of neighborhood size.

Neighborhood Size	Number of Neighborhoods
5	102
10	90
15	78
20	71

Setting  $n_{ki} = 10$  resulted in 90 MSS, and this result is used in subsequent analyses. Given n = 150, K = 90,  $n_k = 10$ , on average sample units belonged to six neighborhoods. In many cases, the similarity of the second element to join the set is low, but the monotonic decline in similarity of subsequent elements is relatively gentle (Figure 2); the lowest similarity of any member of a set ranges from 0.067 to 0.50 with a median of 0.35.

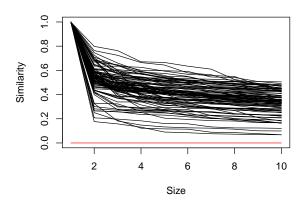


Figure 2. Trace of similarity values for each of 90 MSS as new elements are added to the sets.

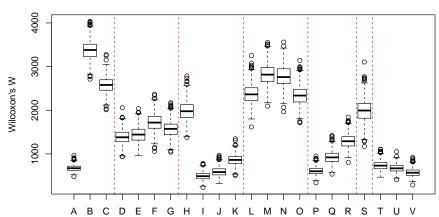
### 3.2. Analysis of Environmental Constraints

Figure 3 shows the distribution of Wilcoxon's *W* for a broad range of interval- or ratio-scaled environmental attributes, mostly related to topography and climate. Boxes A – C represent commonly

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observed plot-level observations: elevation (m), aspect value ( $(\cos(aspect - 30^\circ) + 1)/2$ ), and slope steepness in percent. Elevation shows quite strong constraint, where sample units with similar vegetation composition must occur within a relatively narrow band of elevation; aspect and slope show very little constraint. Boxes D – G are seasonal precipitation and show moderate constraint on similarity. Boxes H – K show seasonal temperature and except for winter temperature (H) show quite significant constraint; spring temperature has the strongest constraint of any attribute, and summer temperature ranks third in effect size. Boxes L – O show direct solar radiation (heat load) and exhibit little direct constraint. Boxes P – R show seasonal potential evapotranspiration (PET); spring PET ranks fourth in effect size, and summer and autumn PET rank ninth and tenth respectively. Boxes T – V show annual climatic summaries: mean annual temperature, number of frost free days, and growing degree days (sum of temperature > 5°C). All three variables show strong constraint; growing degree days ranks second in effect size, the number of frost free days ranks fifth, and mean annual temperature ranks seventh.

# Maximally Similar Sets of Size 10



**Figure 3.** Distribution of Wilcoxon's W for selected environmental attributes with neighborhood size set to 10. A = elevation (m), B = aspect value, C = slope steepness, D = winter precipitation (mm), E = spring precipitation (mm), F = summer precipitation (mm), G = autumn precipitation (mm), H = winter temperature (°C), I = spring temperature (°C), J = summer temperature (°C), K = autumn temperature (°C), L = winter radiation ( $W/m^2$ ), M = spring radiation ( $W/m^2$ ), N = summer radiation ( $W/m^2$ ), O = autumn radiation ( $W/m^2$ ), P = spring potential evapotranspiration (mm), Q = summer potential evapotranspiration (mm), R = autumn potential evapotranspiration, S = site water balance (precipitation - potential evapotranspiration), T = mean annual temperature (°C), U = frost free days, V = growing degree days. Red dashed lines separate logically related attributes.

Figure 4 shows the distribution of Wilcoxon's W for a range of categorical variables, mostly related to soil properties and geology. Boxes A-C show the constraints associated with typically observed soil properties. Of the three, soil depth (classified into four categories) shows the greatest constraint. Boxes D-F show the effect size of soil classifications commonly employed in the United States. Soil short family (D) was classified into 58 classes with many singletons and one class with 21 sample units. Soil subgroup was classified into 16 classes with a few singletons and one class with 50 sample units. Soil great group (F) showed the greatest constraint of the three, and was classified into seven classes with two singletons and maximum of 63 sample units/class.



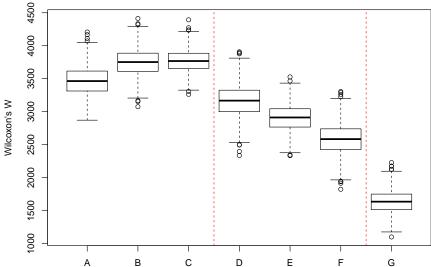


Figure 4. Distribution of Wilcoxon's W for selected environmental attributes with neighborhood size set to 10. A = soil depth, B = soil texture, C = coarse vs fine texture, D = soil short family, E = soil subgroup, F = soil great group, and G = surficial geology. Red dashed lines separate logically related attributes.

Surficial geology was classified into 23 classes, with several singletons and a maximum or 30 sample units/class, and showed the greatest constraint of any of the categorical attributes. Sample units with similar vegetation are likely to occur on a narrow range of surficial geology types.

In general, among the interval-scaled attributes the greatest constraint was demonstrated by variables associated with sample unit temperatures, either directly or indirectly. Spring potential evapotranspiration also ranked in the top five, but is also a function of temperature. Surprisingly, precipitation generally showed little effect, with seasonal precipitation ranking 11-14. Solar radiation showed very little effect, suggesting that the same level of radiation can occur in sites of quite different vegetation composition if the temperature of the units is different. Among the categorical variables only surficial geology showed a substantial constraint, exhibiting values similar to seasonal precipitation among the interval-scale variables.

## 4. Discussion

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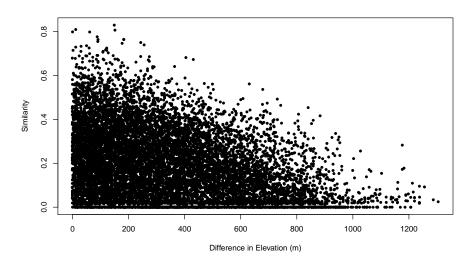
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## 4.1. Constraint vs Determinant

Throughout the results I have repeatedly specified "constraint" as opposed to "determinant" in interpreting the results. The composition of ecological communities (forest vegetation in this case) is determined by a complex set of processes. While species have individual responses to specific environmental attributes, the form of integration of those individual responses into the overall species response is generally unknown. Often, environmental attributes can be partly compensatory, and the relationships are generally not linear or independent. Notably, suitable habitat is necessary but not sufficient, so that species may be absent from a community for reasons unrelated to the environment at that location. Accordingly, the relationship between environment and community composition is a relation, as opposed to a function.

Figure 5 shows the relationship between the pair-wise compositional similarities and the pair-wise differences in sample unit elevation. The plot shows the classical triangular distribution



**Figure 5.** Distribution of sample unit pair-wise similarity as a function of sample unit pair-wise difference in elevation (m).

characteristic of these relationships. As the difference in elevation between sample units increases it becomes increasingly difficult to compensate for that difference, and maximum possible sample unit similarity declines. However, even at zero or small differences in elevation compositional similarity can be zero or low. Given the triangular distribution of the similarity/environment relation, the MSS analysis proposed here looks for boundary conditions, i.e. "how different can values of an environmental attribute be and still allow similar community composition?" While it may appear that such relations are suitable for analysis by logistic quantile regression [5] it's important to note that the points on the figure are not independent of each other, and that in fact every point is associated with N-1 other points because they pertain to the same sample unit.

## 4.2. Neighborhood Size

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The calculation of maximally similar sets was conducted at sizes of 5, 10, 15, and 20. Specific results were only shown for sets of size 10. Across the range of neighborhood sizes the results were quite similar (Spearman rank correlations in the range [0.9282656, 0.9910036]. Nonetheless, neighborhood size does matter in the analysis. Since we are looking for boundary conditions, the maximum pair-wise difference in a neighborhood (and thus statistical power) scales as  $(n_k^2 - n_k)/2$ . Accordingly, small neighborhoods may not exhibit any sample unit pairs at the limit for a given environmental attribute, and thus larger neighborhoods are preferred. However, as neighborhood size increases, the similarity to the neighborhood of late-joining sample units may be quite low, and thus permit quite large pair-wise differences in environment to obtain. The optimal neighborhood size is thus a function of the size of the data set and the distribution of similarities in the similarity relation, and generally cannot be known a priori.

## 4.3. Interval-Scaled vs Categorical Variables

Both interval-valued and categorical variables were analyzed using the Wilcoxon rank sum test, and thus present results on the same scale. However, depending on the number of classes within a categorical variable, it's common to get observed or random sets that have identical entropies, and thus generate ties in the calculation of Wilcoxon's W. As the number of ties goes up, the power of the

test declines, and some categorical variables thus have low power. In addition, ideally the distribution of cases by class would be relatively balanced. In the data analyzed here the distributions were wildly skewed, which again reduces power. Of the categorical variables analyzed, only surficial geology demonstrated strong constraint. The extent to which the other variables suffered from low power, as opposed to minimal ecological effect, is difficult to know, but the results obtained do make sense from an ecological perspective.

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#### 195 Abbreviations

196 The following abbreviations are used in this manuscript:

MSS Maximally Similar Sets
PET potential evapotranspiration

#### 199 References

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#### 209 Appendix A.

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The Shoshone National Forest in Wyoming, USA, is part of the United States Department of Agriculture National Forest System. In a previous analysis of plant community composition 1204 sample units of 375 m<sup>2</sup> were stratified throughout the National Forest. The total abundance of all individuals of each vascular plant species was estimated as the percent of the plot area covered by foliage of that species recorded in the classes given by Table A1.

Table A1. Species Abundance Classes

Range	Mid-Point Abundance
0 ≤ 1	0.5
$1 \le 5$	3
$5 \le 15$	10
$15 \leq 25$	20
$25 \le 35$	30
$35 \leq 45$	40
$45 \le 55$	50
$55 \le 65$	60
$65 \le 75$	70
$75 \le 85$	80

Of the original 1204 sample units, 150 sample units of forest communities were drawn at random from the pool for the analysis presented here. The natural log of the mid-point abundances given in Table A1 were used in the calculation of the sample unit similarity relation as given in equation 4.

Environmental variables were either recorded in the field, or modeled in a GIS. Field measured variables are in table A2.

Table A2. Field Measured Variables

Variable	Description
elevation	elevation in meters above sea level
aspect value	$(\cos(aspect-30)+1)/2$
slope	slope gradient in percent
soil depth	shallow, moderate, deep, or very deep
parent material class	coarse loamy, fine loamy, loamy skeletal, sandy skeletal, or fragmented
texture	coarse or fine
short family	USDA Soils classification (58 classes)
soil subgroup	USDA Soils classification (16 classes)
soil great group	USDA Soils classification (7 classes)
surficial geology	geologic rock type (23 classes)

## GIS modeled data are in Table A3.

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Table A3. GIS Modeled Variables

Variable	Description
Winter	January, February, and March
	sum of precipitation (mm)
	mean monthly temperature (°C)
	solar radiation $(W/m^2)$
Spring	April, May and June
	sum of precipitation (cm)
	mean monthly temperature (°C)
	solar radiation $(W/m^2)$
	potential evapotranspiration
Summer	July, August, September
	sum of precipitation (cm)
	mean monthly temperature (°C)
	solar radiation $(W/m^2)$
	potential evapotranspiration
Autumn	October, November, and December
	sum of precipitation (cm)
	mean monthly temperature (°C)
	solar radiation $(W/m^2)$
	potential evapotranspiration
Frost-free days	length of growing season in days
degree days	sum of hours $> 5^{\circ}$ C
site water balance	sum of (precipitation - potential evapotranspiration)
mean annual temperature	°C

The sample unit vegetation and site data are available at Figshare at https://doi.org/10.6084/m9.figshare.7234121.v1

The R code for the analysis is available at Figshare at https://doi.org/10.6084/m9.figshare.7234124.v1