

PRIMER / PERMANOVA Essentials

Outline of Topics

- 1. Properties of multivariate data
 - a. Pre-treatment options, including transformation, standardization, normalization & dispersion weighting
 - b. Distance/dissimilarity/similarity measures for a variety of data types
- 2. Cluster analysis (CLUSTER), including tests for significant structure within clusters (SIMPROF) to permit non-arbitrary classifications of samples or variables (species)
- 3. Ordination via projection:
 - a. Principal component analysis (PCA); and
 - b. Principal coordinate analysis (PCO)
- 4. Ordination to preserve inter-sample relationships via non-metric, metric or threshold metric multi-dimensional scaling (nMDS, mMDS, tmMDS)
- 5. Relating biotic to abiotic data, including tests of association between resemblance matrices (RELATE)
- 6. Finding optimal subsets of environmental (or other) variables that generate a 'best' match to patterns among samples based on species variables (BEST BIOENV)
- 7. Non-parametric permutation tests for differences among a priori groups of samples:
 - a. Analysis of Similarities (ANOSIM);
 - b. Ordination of Bootstrap averages; and
 - c. Finding important variables (SIMPER and BEST BVSTEP)
- 8. Diversity measures; Taxonomic distinctness; Taxonomic resemblance
- 9. Partitioning variation for high-dimensional data on the basis of a resemblance measure of choice and tests for differences in centroids (location) for a priori groups (PERMANOVA), including:
 - a. One-way and two-way cases & constructing a.priori contrasts;
 - b. Tests for interactions; and
 - c. Pairwise tests
- 10. Multivariate variation (spread), tests for homogeneity of multivariate dispersions and comparisons of beta diversity (PERMDISP); PERMANOVA tests for centroid differences in the presence of heterogeneity.
- 11. Complex multi-factor experimental designs (PERMANOVA), including:
 - a. Fixed/random factors; Crossed/nested relationships; Expectations of mean squares; Components of variation; Types of SS; Covariates; Pooling; Unbalanced designs
 - b. Centroid plots for Main Effects and Interactions; Residual plots.
- 12. Fitting multivariate response data (e.g., species) to continuous predictor variables (e.g., environmental), including:
 - a. model selection (DISTLM); and
 - b. dissimilarity-based redundancy analysis (dbRDA) to visualise fitted variation
- 13. Identifying when an observation is 'unusual', given previous observations (e.g., in a time-series / monitoring program) using multivariate dissimilarity-based Control Charts.
- 14. Wrap-up and Overview of the week / Open Q&A Session / Own-data Session.

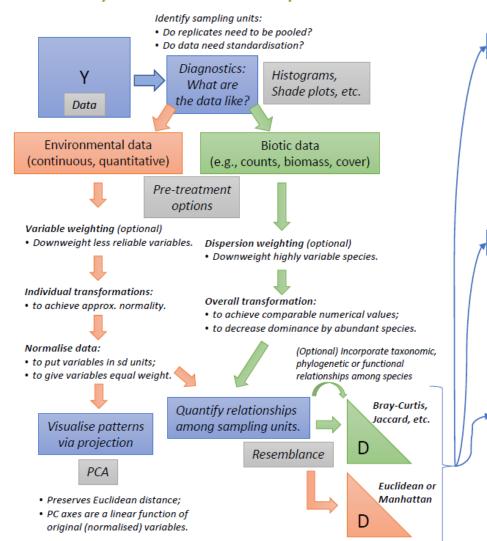
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Provisional Time-Table

The time-table below should be taken as a rough guide. Breaks will be taken at the specified times each day, but individual topics may flow over or under allotted time-slots, depending on the depth of coverage, questions asked and ensuing discussions. Participants will be given real data sets to analyse during practical sessions. Participants are encouraged to ask questions and discuss their own data with the presenter throughout the week.

	Monday	Tuesday	Wednesday	Thursday	Friday
Session 1 8:30 – 10:30	1. Pre-treatment options; resemblance measures	4. Ordination with nMDS; mMDS	7a,b. ANOSIM; Bootstrap averages	11. Unbalanced case; Test centroids when dispersions differ (PERMANOVA)	12. DISTLM / dbRDA
Coffee Break 10:30 – 11:00					
Session 2 11:00 – 12:30	2. CLUSTER; SIMPROF (cont'd)	4. Ordination with mMDS; mMDS (cont'd)	7c. SIMPER; BVSTEP; 8. Diversity	11. Fixed/random; crossed/nested; (PERMANOVA)	13. Control Charts
Lunch 12:30 – 13:30					
Session 3 13:30 – 15:30	3a. Ordination with PCA	5. RELATE; seriation or cyclical models	9. PERMANOVA: one-way; two-way; interactions; pairwise tests; contrasts	11. Complex designs; mixed models; pooling (PERMANOVA)	14. Wrap up / Overview / Q&A
Coffee Break 15:30 – 16:00					
Session 4 16:00 – 17:30	3b. Ordination with PCO	6. BEST; BIOENV; global test	10. PERMDISP; beta diversity	11. Finite factors; asymmetrical designs (PERMANOVA)	14. Q&A / 'Own- data' Session

Analysis Pathways in PRIMER



Cluster analysis

Identify associations/groups of samples or species.

- CLUSTER hierarchical agglomerative
- UNCTREE binary divisive
- · LINKTREE constrain binary splits by env vars.
- kRCLUSTER maximise ANOSIM R for given k

SIMPROF

Identify "significant" associations among sampling units or species.

Ordination

Visualise patterns of relationships/ Reduce dimensionality

- mMDS metric MDS
- tmMDS threshold-metric MDS
- nMDS non-metric MDS

Bootstrap averages

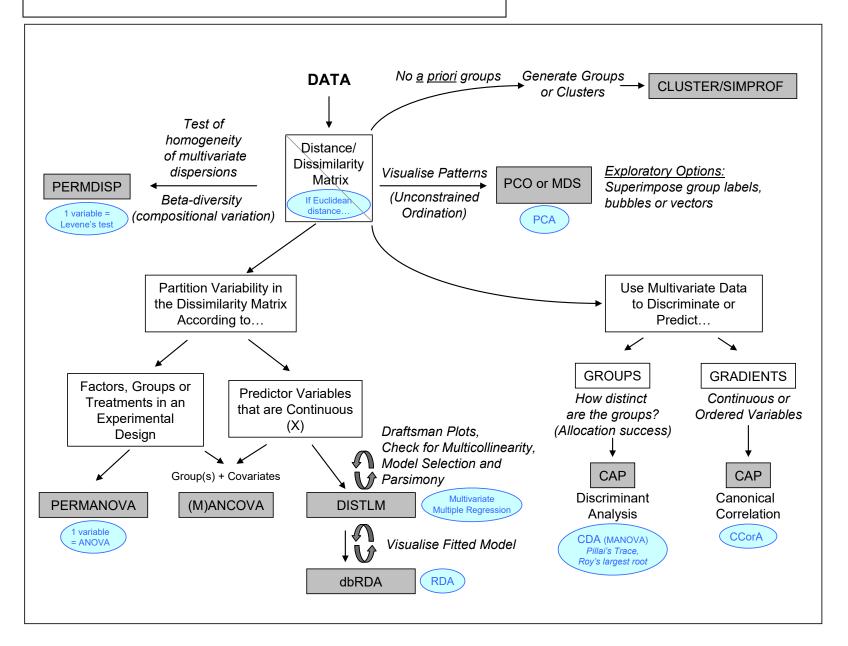
Visualise average values and their variability in reduced-dimension plot.

Hypothesis-testing

Robust, non-parametric permutation-based p-values.

- · ANOSIM compare groups (ordered or unordered), R
- RELATE relate two resemblance matrices (rho).
- BIOENV find subsets of vars that maximise rho.

Overview of Tools and Methods in PERMANOVA+



Some key and well-cited papers on PRIMER / PERMANOVA+ methods

PRIMER

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- Clarke KR (1993) Non-parametric multivariate analyses of changes in community structure. *Aust J Ecol* 18: 117-143
- Clarke KR (1999) Non-metric multivariate analysis in community-level ecotoxicology. *Environ Toxicol Chem* 18: 118-127
- Clarke KR, Ainsworth M (1993) A method of linking multivariate community structure to environmental variables. *Mar Ecol Prog Ser* 92: 205-219
- Clarke KR, Chapman MG, Somerfield PJ, Needham HR (2006) Dispersion-based weighting of species counts in assemblage analyses. *Mar Ecol Prog Ser* 320: 11-27
- Clarke KR, Gorley RN (2001, 2006, 2015) *PRIMER v5, v6, v7: User manual/tutorial.* PRIMER-E, Plymouth, UK, 91pp, 192pp, 296pp
- Clarke KR, Green RH (1988) Statistical design and analysis for a 'biological effects' study. *Mar Ecol Prog Ser* 46: 213-226
- Clarke KR, Somerfield PJ, Airoldi L, Warwick RM (2006) Exploring interactions by second-stage community analyses. *J Exp Mar Biol Ecol* 338: 179-192
- Clarke KR, Somerfield PJ, Chapman MG (2006) On resemblance measures for ecological studies, including taxonomic dissimilarities and a zero-adjusted Bray-Curtis coefficient for denuded assemblages. *J Exp Mar Biol Ecol* 330: 55-80
- Clarke KR, Somerfield PJ, Gorley RN (2008). Exploratory null hypothesis testing for community data: similarity profiles and biota-environment linkage. *J Exp Mar Biol Ecol* 366: 56-69
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- Clarke KR, Tweedley JR, Valesini FJ (2014) Simple shade plots aid better long-term choices of data pretreatment in multivariate assemblage studies. *J Mar Biol Assoc UK* 94: 1-16
- Clarke KR, Warwick RM (1994, 2001, 2014) Change in Marine Communities: An Approach to Statistical Analysis and Interpretation. PRIMER-E, Plymouth, UK. 1st ed: 144pp; 2nd ed: 172pp. 3rd ed: (authors: Clarke KR, Gorley RN, Somerfield PJ, Warwick RM) 260pp
- Clarke KR, Warwick RM (1998) Quantifying structural redundancy in ecological communities. *Oecologia* 113: 278-289
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- Clarke KR, Warwick RM (2001) A further biodiversity index applicable to species lists: variation in taxonomic distinctness. *Mar Ecol Prog Ser* 216: 265-278
- Field JG, Clarke KR, Warwick RM (1982) A practical strategy for analysing multispecies distribution patterns. *Mar Ecol Prog Ser* 8: 37-52
- Somerfield PJ, Clarke KR (1995) Taxonomic levels, in marine community studies, revisited. *Mar Ecol Prog Ser* 127: 113-119
- Somerfield PJ, Clarke KR (2013) Inverse analysis in non-parametric multivariate analyses: distinguishing groups of associated species which covary coherently across samples. *J Exp Mar Biol Ecol* 449: 261-273
- Somerfield PJ, Clarke KR, Gorley RN (2021a) A generalized analysis of similarities (ANOSIM) statistic for designs with ordered factors. *Austral Ecology* 46: 901-910.
- Somerfield PJ, Clarke KR, Gorley RN (2021b) Analysis of similarities (ANOSIM) for 2-way layouts using a generalized ANOSIM statistic, with comparative notes on Permutational Multivariate Analysis of Variance (PERMANOVA). *Austral Ecology* 46: 911-926.
- Somerfield PJ, Clarke KR, Gorley RN (2021c) Analysis of similarities (ANOSIM) for 3-way designs. *Austral Ecology* 46: 927-941.

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- Warwick RM, Clarke KR (1991) A comparison of some methods for analysing changes in benthic community structure. *J Mar Biol Ass UK* 71: 225-244
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PERMANOVA+

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- Anderson MJ (2001) Permutation tests for univariate or multivariate analysis of variance and regression. Can J Fish Aquat Sci 58: 626-639
- Anderson MJ (2006) Distance-based tests for homogeneity of multivariate dispersions. *Biometrics* 62: 245-253
- Anderson MJ (2008) Animal-sediment relationships revisited: characterising species' distributions along an environmental gradient using canonical analysis and quantile regression splines. *J Exp Mar Biol Ecol* 366: 16-27
- Anderson MJ (2017) Permutational Multivariate Analysis of Variance (PERMANOVA). Wiley StatsRef: Statistics Reference Online. 1 15. Article ID: stat07841. https://doi.org/10.1002/9781118445112.stat07841
- Anderson MJ, Crist TO, Chase JM, Vellend M, Inouye BD, Freestone AL, Sanders NJ, Cornell HV, Comita LS, Davies KF, Harrison SP, Kraft NJB, Stegen JC, Swenson NG (2011) Navigating the multiple meanings of β diversity: a roadmap for the practicing ecologist. *Ecol Lett* 14: 19-28
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- Anderson MJ, Legendre P (1999) An empirical comparison of permutation methods for tests of partial regression coefficients in a linear model. *J Statist Comput Sim* 62: 271-303
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- Anderson MJ, Robinson J (2003) Generalized discriminant analysis based on distances. *Aust NZ J Stat* 45: 301-318
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