Bio3018F Practical 2022

Biodiversity and Ecosystem Function in the Cape Floristic Region

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Introduction

Our understanding of and approach to studying the relationship between biodiversity and ecosystem function (B-EF) has evolved over the past few decades (Figure 1; van der Plas 2019). The historical view was one of biodiversity as the response variable, being determined by environmental and anthropogenic factors, with little feedback to ecosystem function. In the early 1990s, this shifted (and perhaps overcompensated) to focus on the causal effects of variation in biodiversity on ecosystem functioning, with little emphasis on the role of environmental variation. More recently, there is recognition that biodiversity both responds to the environment and partly drives ecosystem function in concert with environmental variation. The current focus of most B-EF research is on the relative importance of abiotic drivers (natural and anthropogenic) versus biotic variation in determining various ecosystem functions.

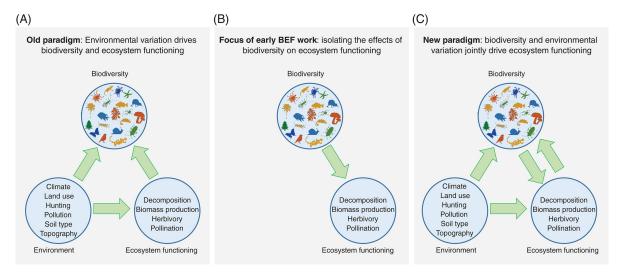


Figure 1: (from van der Plas 2019)

The Cape Floristic Region (CFR) of South Africa is one of the most botanically diverse areas on the planet. The indigenous flora of the CFR has several components with different evolutionary and biogeographic origins (Figure 2; Bergh et al. 2014), and distinct differences in a range of ecosystem functions. The CFR also has a long history of global change impacts, from direct anthropogenic disturbance (e.g. land use / land cover) to the introduction of invasive alien species. This provides a range of highly varied ecosystem types within close proximity, that often share or contrast in their biotic composition (species, functional and phylogenetic diversity) and abiotic conditions - climate, soils and disturbance regimes (fire, herbivory).

[Describe the measures of ecosystem function here... i.e. average and amplitude of seasonality of the Normalized Difference Vegetation Index (NDVI) recorded by the MODIS Terra satellite...]

In this practical, we will explore how variation in a set of measures of biodiversity (species, functional and

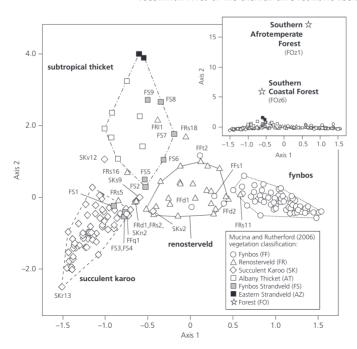


Figure 2: Ordination of genus-level floristic similarities of GCFR vegetation units sensu Mucina and Rutherford (2006), as inferred on the basis of the 'important species' lists provided in The vegetation of South Africa, Lesotho and Swaziland (Mucina and Rutherford 2006). Figure from Bergh et al. 2014.

phylogenetic α and β diversity) and environmental conditions relate to a set of metrics of ecosystem function derived from satellite timeseries.

The questions are:

- Does the variation in environmental conditions potentially explain the observed variation in biodiversity (species, functional and phylogenetic α and β diversity)?
- To what degree does the variation in functional and phylogenetic diversity potentially explain the observed variation in our measures of ecosystem function?
- What is the relative role of the environment versus biodiversity in determining the observed variation in ecosystem function?

Finally, consider this practical a descriptive study. In your discussion, describe a follow-up study that you would perform to discern cause from correlation and partition the relative influence of environmental conditions versus biodiversity on our measures of ecosystem function.

Methods:

The sites are selected to represent contrasting vegetation, but to all be the same (or similar) post-fire age (time since last fire). We will split into 4 teams of 2 or 3. Each pair will survey a point location (towards the corners) within the 250 by 250m MODIS pixel that makes up each site.

Each team will need:

- One or more smartphones
- 2 x 10m measuring tape (or longer)
- 1 x clipboard, paper and pen or pencil

- 1 x densiometer
- 1 x metre rule
- 1 x Vernier calipers
- 8 x large plastic bags for carrying soil and plant samples for each point location
- 8 x soil sample bags
- 2 x masking tape
- 2 x marker pen

At each site, navigate to your team's point location, lay out the two tapes at 90 degrees, crossing at 3.2m and ending at 6.4m. Consider this your guide for a 3.2m radius (~20m2) circle. Within the circle we will measure:

- 1. Environmental conditions:
- Estimate % projected cover (think the area you would see from above) bare soil
- Take a densiometer reading at ground level. To do this, hold the densiometer level on the soil surface. Pick a spot close to the centre, but try not to pick an obviously open (or closed) patch, it should be representative of the location. To take a reading, split each square into quarters and score them for the amount of light visible a value from 0 (complete canopy cover) to 4 (no vegetation visible) counting up these values for all 24 squares (to a maximum of 96), and write this down. We will convert this to canopy cover later, applying the formula 100 1.04 * X, where X is your reading.
- Take a **soil sample** (as per Mike's prac, but it can be smaller as we are going to pool the 4 samples per site). These we will process for soil colour and pH as per Mike's prac.
- Do a **dung count**, scoring dung for the point location by the number of quarters where you find dung (i.e. a single score from 0-4 for the point location). Try to avoid scoring obvious single scat ("dung creation") events in more than one quadrat, unless it's an impressive pile.
- Take a few notes (and photos) on any other features that you think may be important or interesting (slope, rockiness, whatever).
- 2. Diversity sampling:
- First, make sure your smartphone is fully charged and set to record GPS location with your photographs!
- With your team, decide on which are the top 5 species by % projected cover. If your site is dominated by fewer than 5 species, count up as many species as make up 80% of the vegetation.
 - Take diagnostic photos for the 5 species (habit (whole plant), leaf, base, flowers and/or fruits).
 - For each of three individuals of your 5 target species, measure the height and collect a shoots for measuring leaf length and leaf width (mark with masking tape and put in sample bag for the site).
- Finally, set a timer and take as many photos of new species (other than your 5 target species) within or near your plot as you can before the alarm goes off. Make sure to take a photo of your site label on your sample bag between sites so you know which photos were collected at which sites. When we get back to base you will upload the photos to folders in the intranet labeled by site.

Analyses

Species Diversity

For this I have just taken the count of species encountered at each point, and the aggregated set of unique species for each site. i.e. no rarefaction etc. ou think this is an issue for the method we used?

Site	Species Number
grass	34
invasion	13
renosterveld	28

Site	Species Number
sandstone	24

And by point site:

Species Number
17
12
13
14
13
17
3
12
24

Functional Diversity

Here I've estimated Functional Diversity (FD) according to the method of Petchey and Gaston (2002) for each of the points, sites and functional turnover between points and between sites using the method of Bryant et al. 2008.

Sites:

	FD	SR
invasion	12.181	11
limestone	5.815	11
sandstone	6.964	9
grass	7.369	9
renosterveld	7.483	11
sand	6.876	9

By point sample:

	FD	SR
grass_SE	3.200	2
$renosterveld_SE$	4.181	5
$invasion_SE$	7.422	4
sand_SE	4.650	5
$sandstone_SE$	5.231	5
$limestone_SE$	4.599	5
grass_NE	3.729	3
renosterveld_NE	4.756	4
invasion_NE	5.856	5
sand_NE	4.136	4
sandstone_NE	4.344	3
$limestone_NE$	4.193	5
grass_SW	4.778	5
renosterveld SW	5.410	5
$invasion_SW$	7.150	4

	FD	SR
sand_SW	4.873	4
$limestone_SW$	4.599	5
$sandstone_SW$	4.344	3
$grass_NW$	4.680	3
$renosterveld_NW$	4.018	5
invasion_NW	6.413	3
$\operatorname{sand}_{-}\operatorname{NW}$	2.713	1
$sandstone_NW$	5.840	5
$limestone_NW$	4.193	5

And functional turnover between sites:

	invasion	limestone	sandstone	grass	renosterveld	sand
invasion						
limestone	0.401					
sandstone	0.431	0.487				
grass	0.487	0.481	0.643			
renosterveld	0.378	0.412	0.560	0.603		
sand	0.460	0.504	0.800	0.647	0.564	

See help file ?picante::phylosor in R for details to help know how to interpret. Note that while the function was written for phylogenetic turnover, we've used it for functional turnover.

Phylogenetic Diversity

Here I've estimated Faith's Phylogenetic Diversity (PD; Faith 1991) for each of the sites and phyloSor (Bryant et al. 2008), the phylogenetic turnover between sites.

```
tree <- read.nexus("prac/Soltis_etal_639taxa_8cploci_rooted.dated.tre")
#plot.phylo(tree, type = "radial", cex = 0.25)</pre>
```

Environmental Similarity among sites

Here I provide a table and an ordination of the site-level environmental measurements we took relating to various aspects of soil conditions, canopy coverage and herbivory.

Ecosystem function

To explore ecosystem function we looked at the 20-year time series of the Normalized Difference Vegetation Index (NDVI) recorded by the MODIS satellite mission. From these we used the post-fire recovery trajectory modelling framework developed by Wilson et al (2015) to derive estimates of the mean maximum NDVI (alpha + gamma), and the amplitude and timing of seasonality (big alpha and phi) as our measures of ecosystem function. Here I've provided a table of these parameters by site and plots of the model fits. Are the models good fits? Do they miss anything? Is it relevant to the questions we're asking?

References

Bergh et al

Mucina and Rutherford

Slingsby et al. 2020 van der Plas Wilson et al