

# Studying land snail ecology to inform conservation planning on Norfolk Island



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John Hunt, Kate Umbers

**WESTERN SYDNEY**  
UNIVERSITY



School of Science  
Western Sydney University







# O'ahu, Hawai'i

282 species



# Norfolk Island

60 species







Norfolk Island





Palm



Hardwood



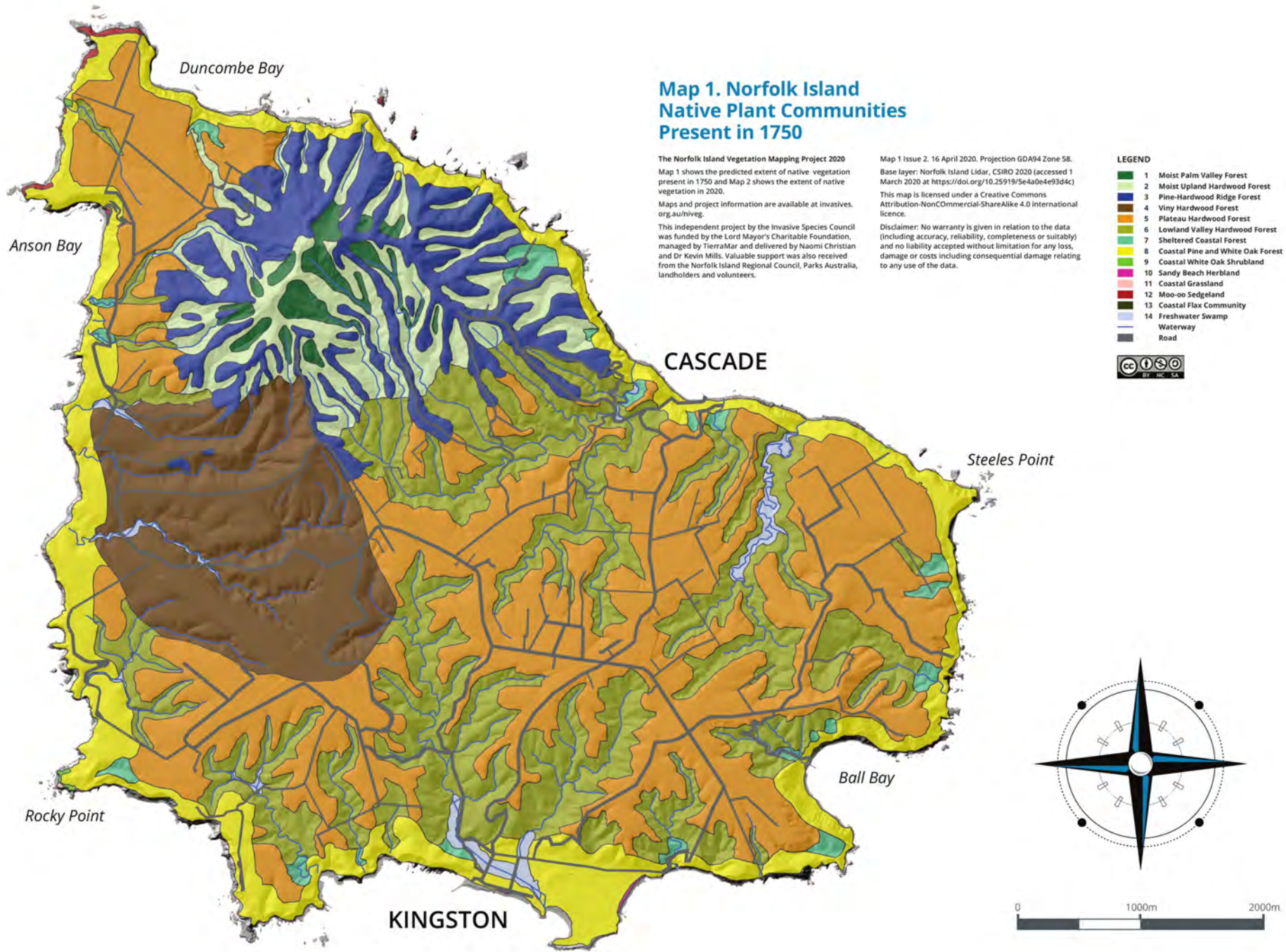
Pine



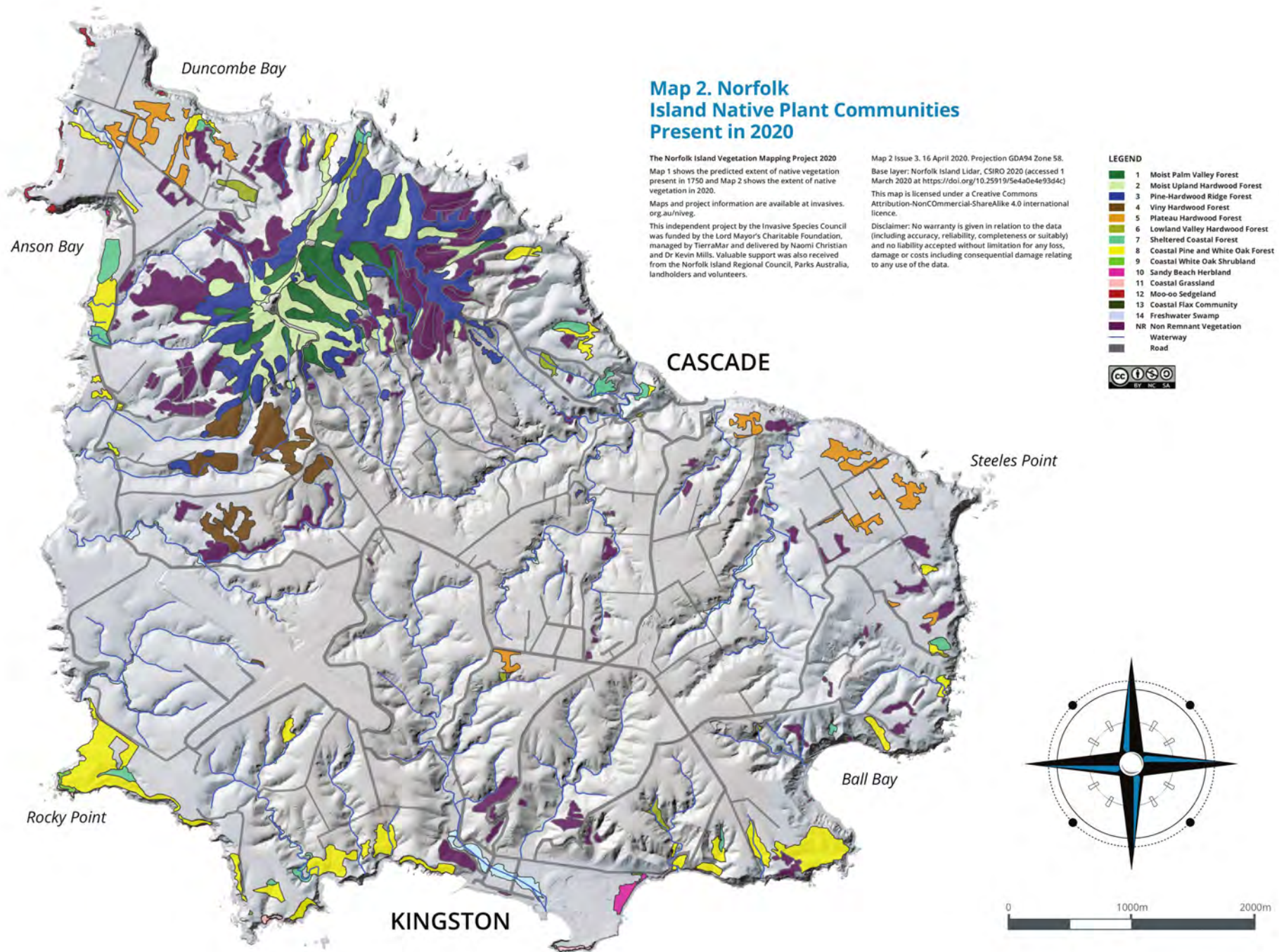
Guava & olive











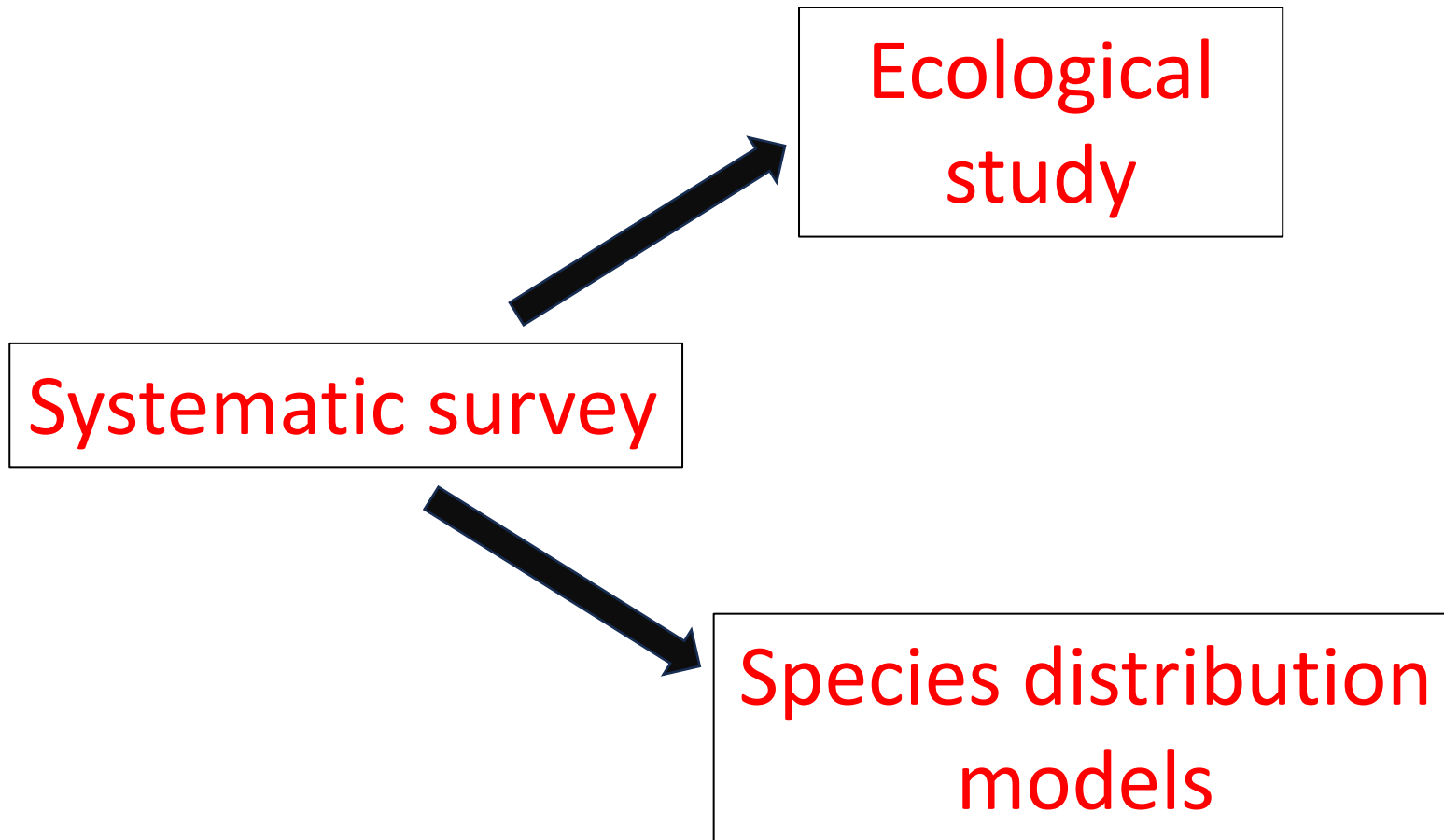


# Land snails of Norfolk Island



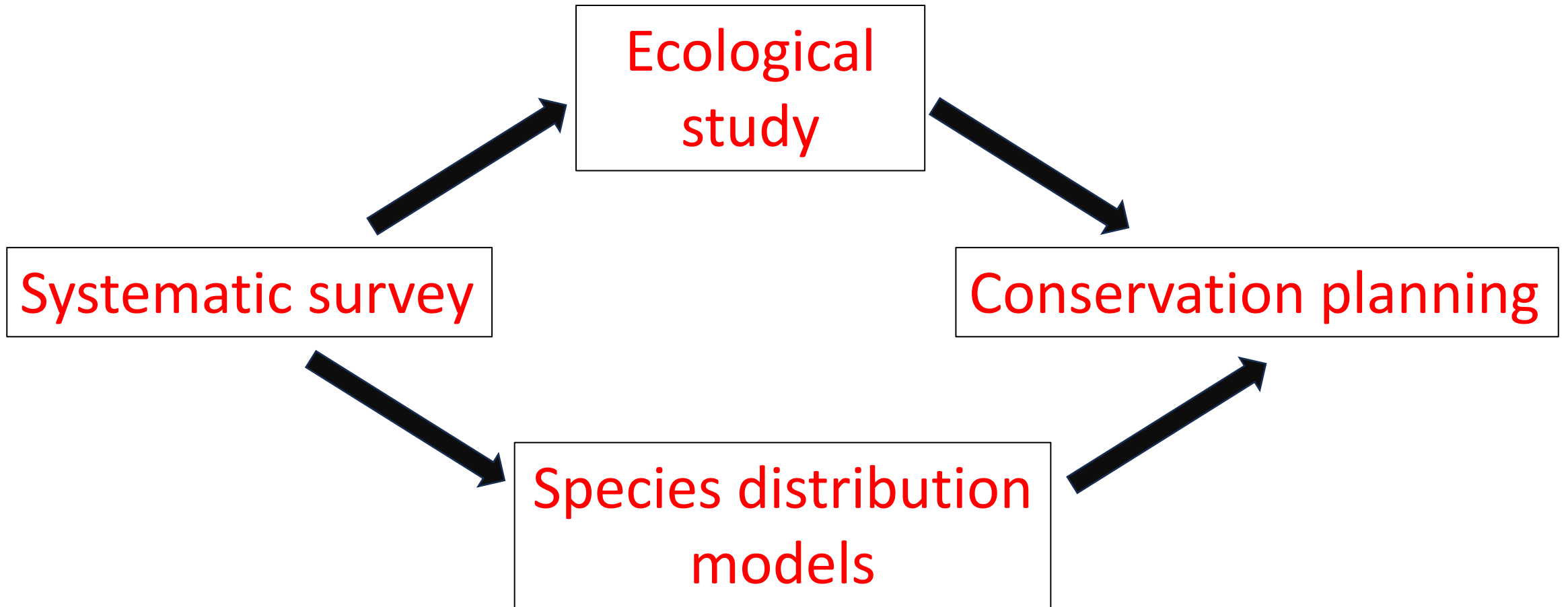


# Key steps





# Key steps

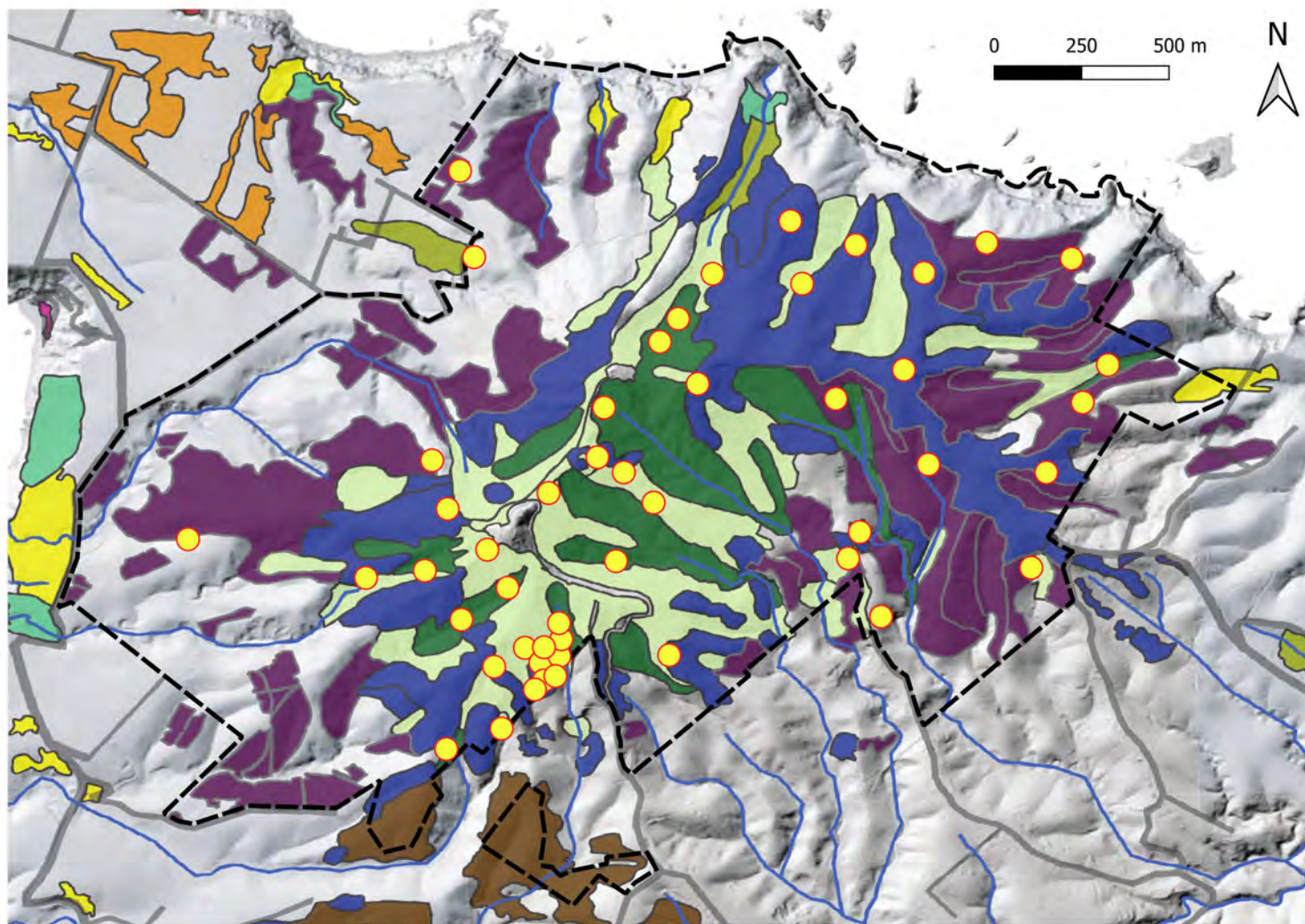




## Question 1

What environmental factors are associated with abundance of a land snail species?





The Norfolk Island Vegetation Mapping Project 2020

## Systematic survey

Sep 2023 – Oct 2023

49 sites

4m x 4m quadrat per site





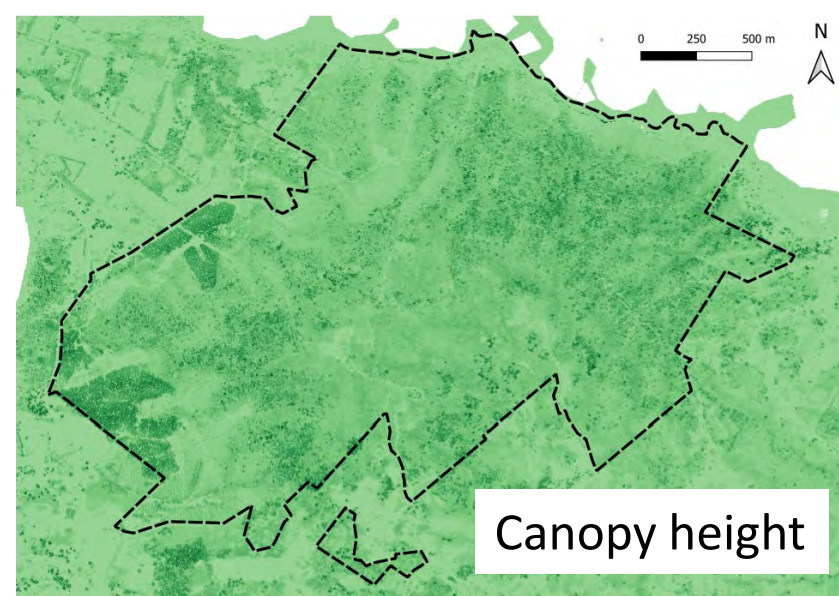
## Data collected

- counts per snail species
- Soil pH
- Elevation
- Plant percentage cover
- Microhabitat percentage cover



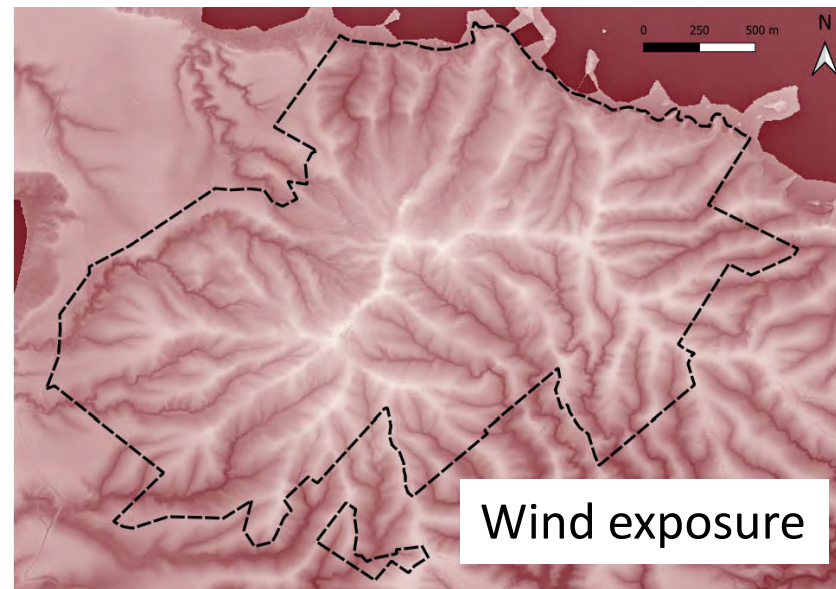
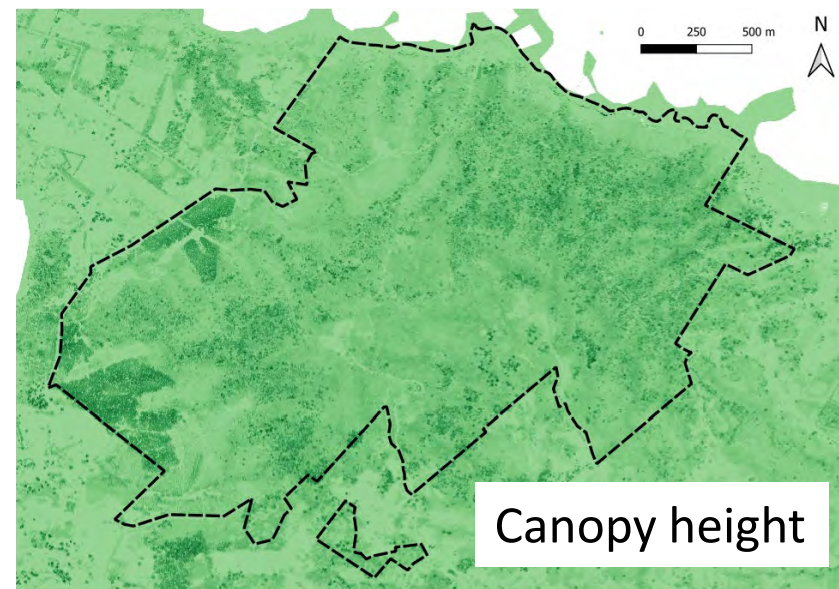


# Environmental variables



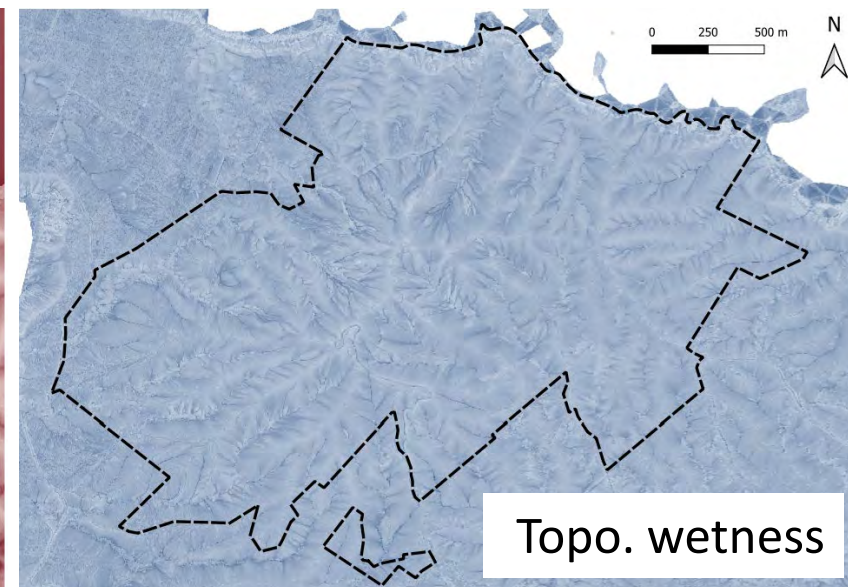
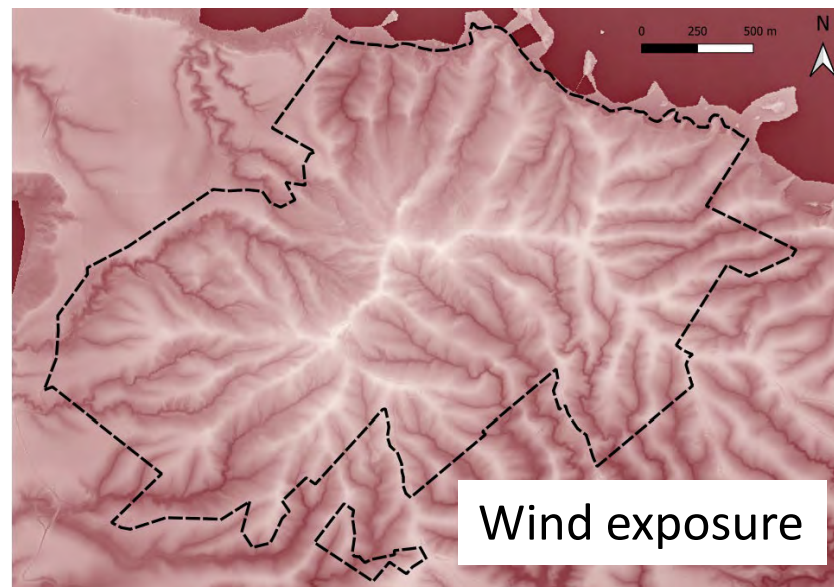
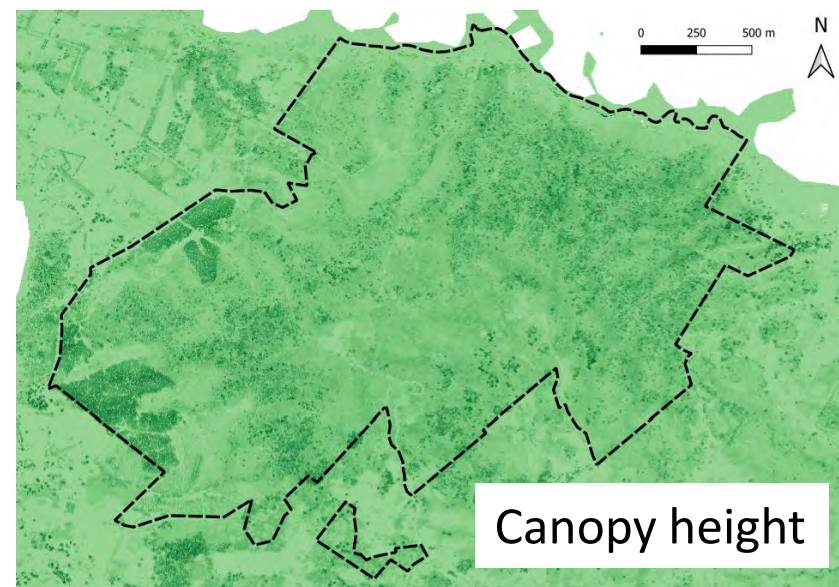


# Environmental variables



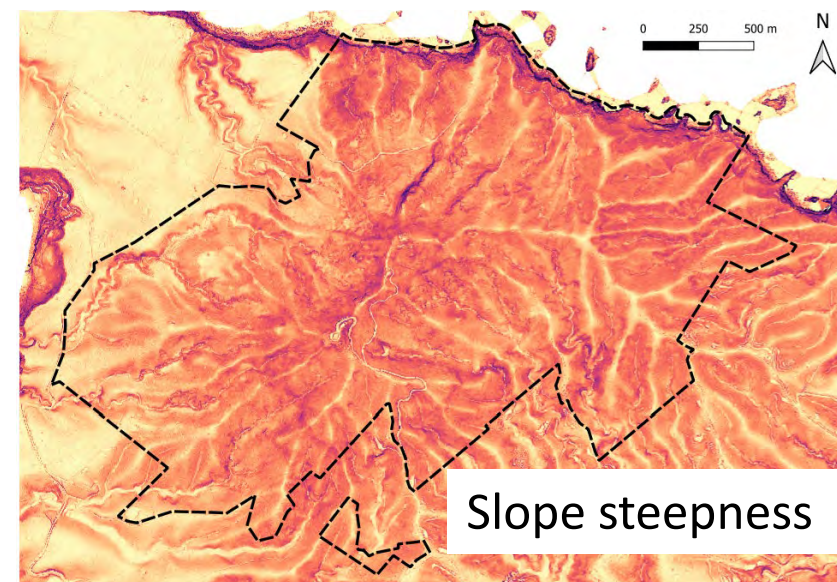
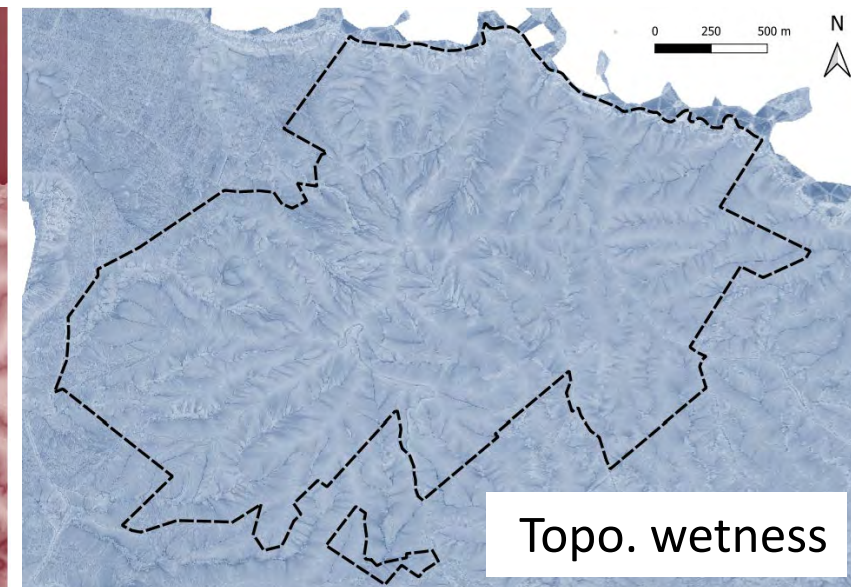
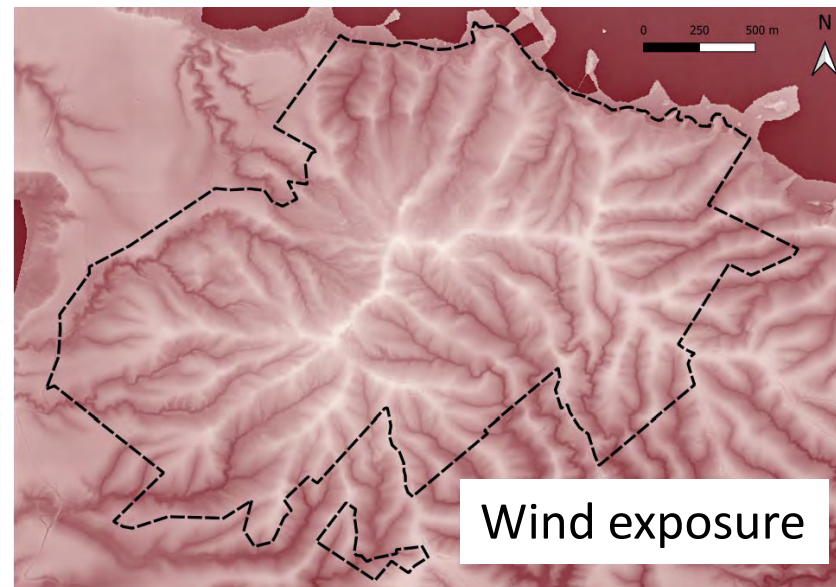
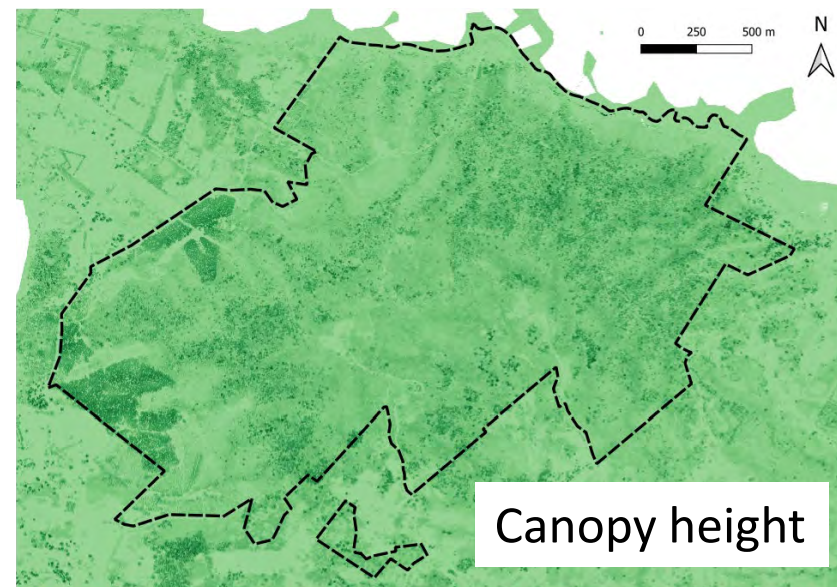


# Environmental variables



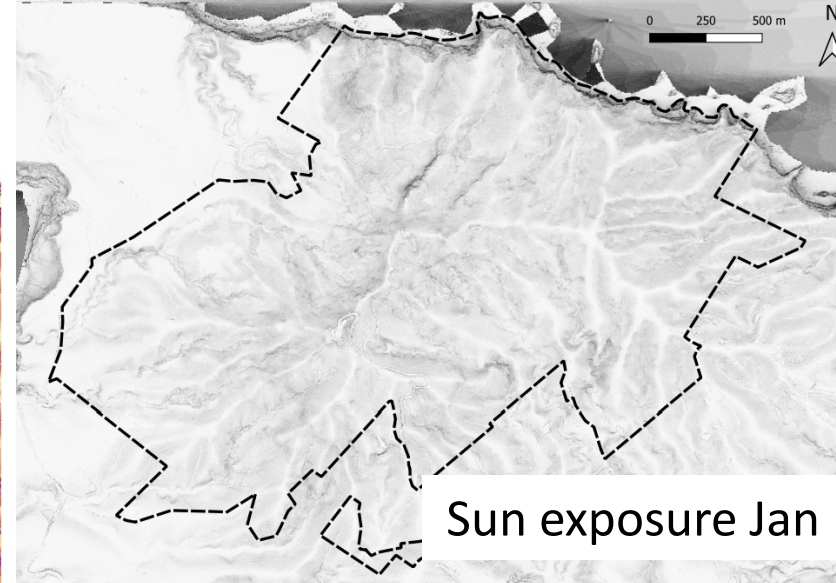
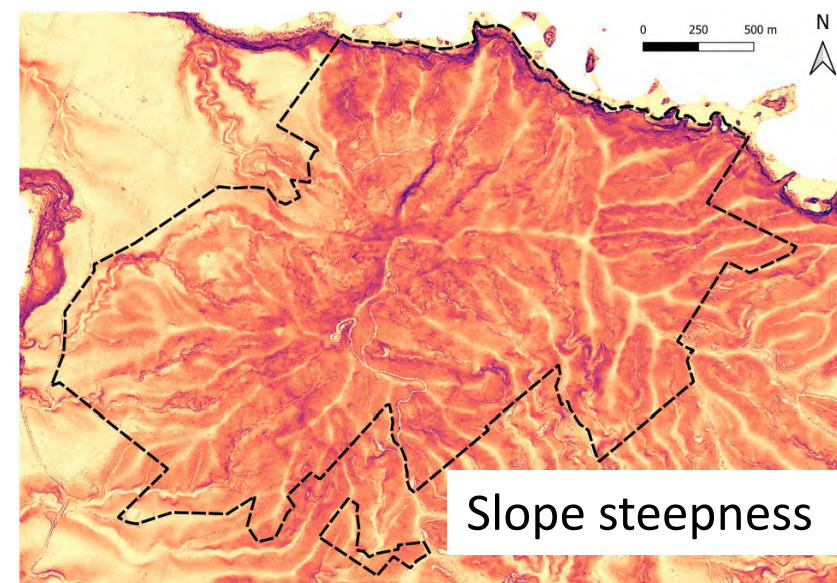
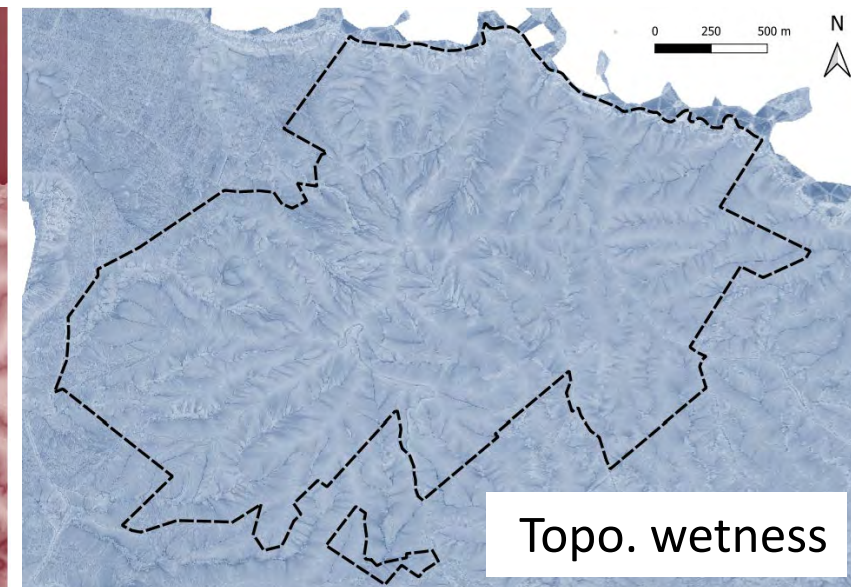
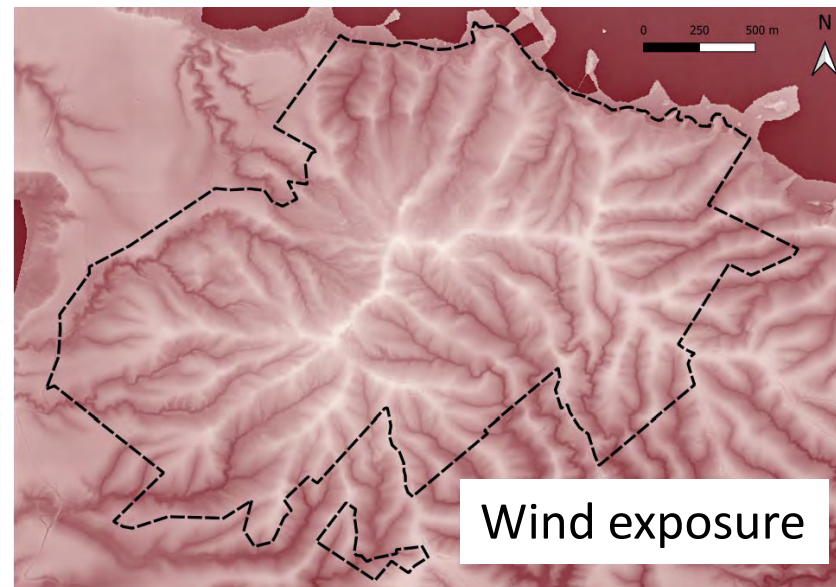
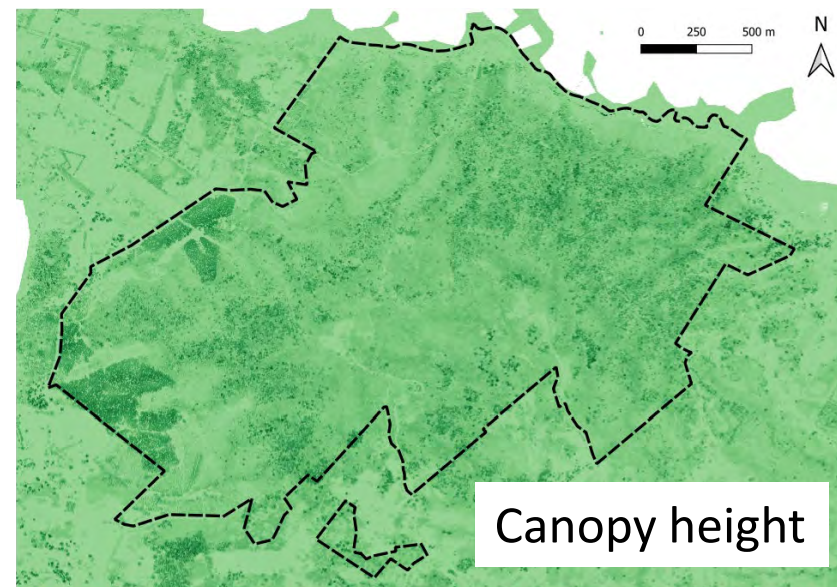


# Environmental variables



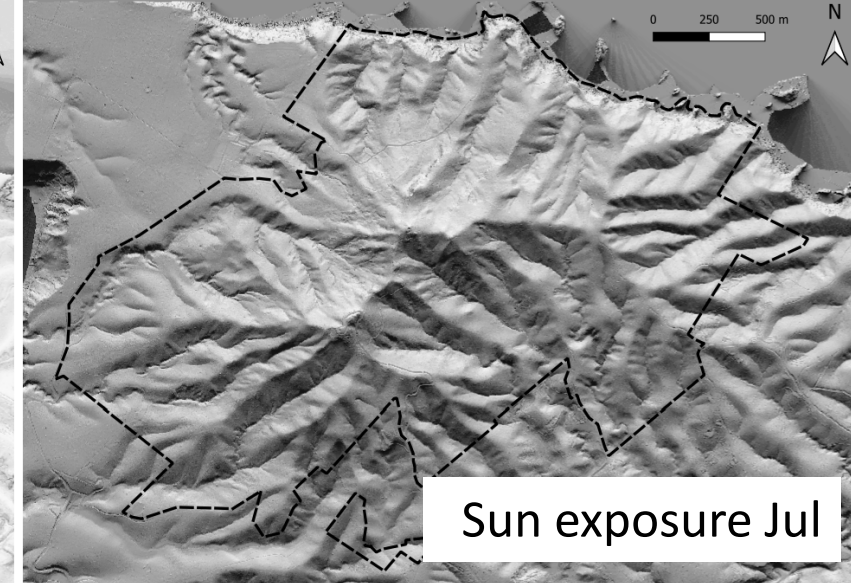
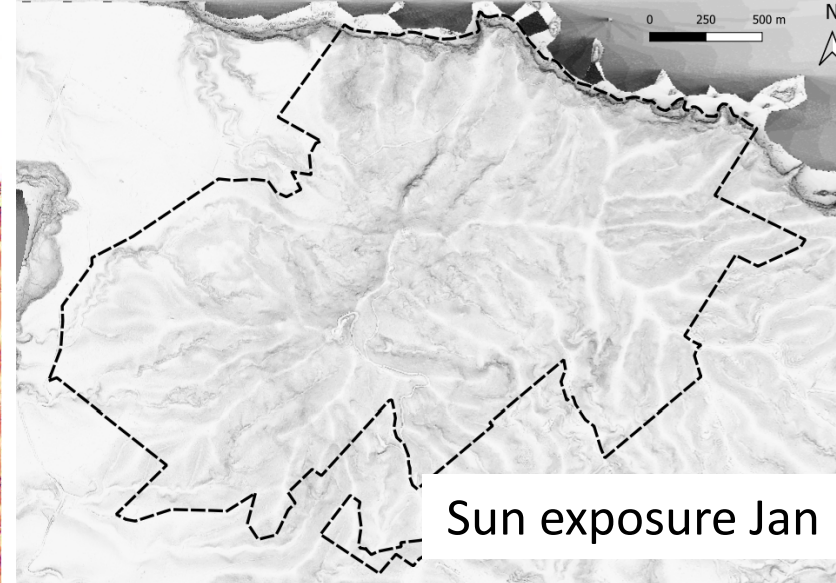
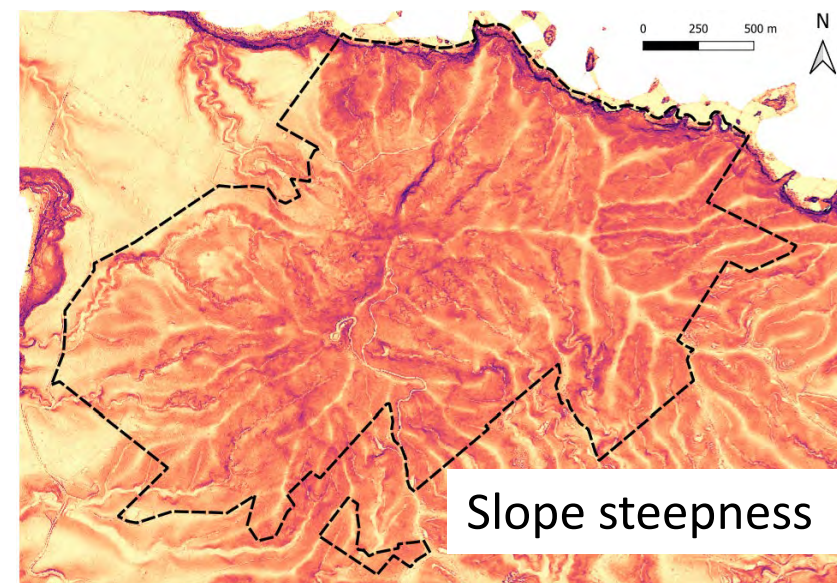
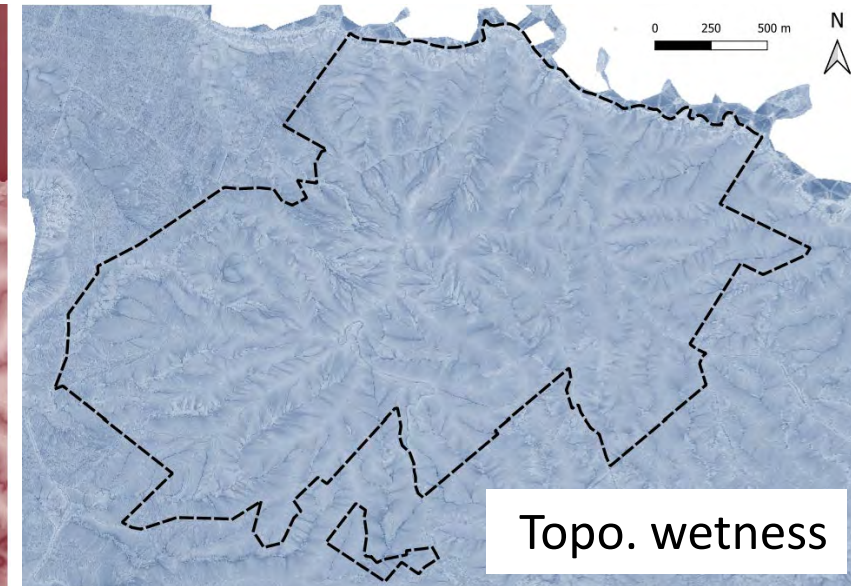
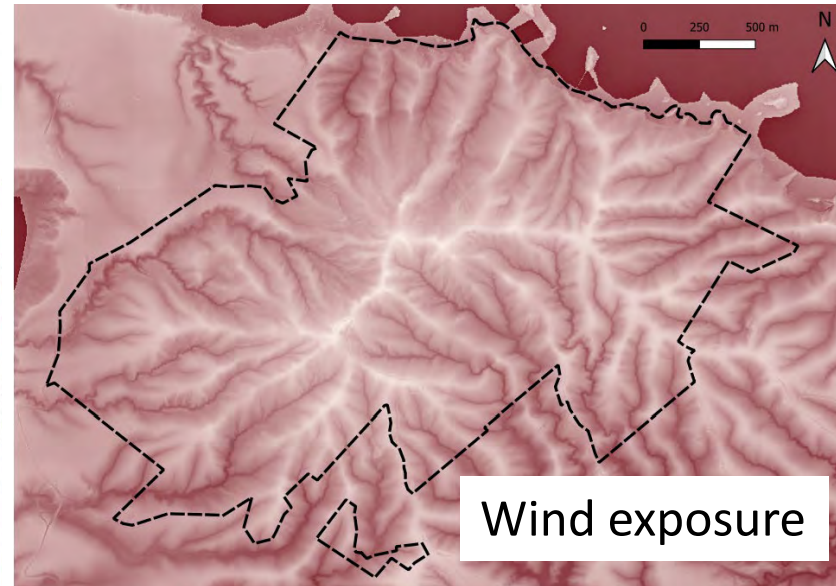
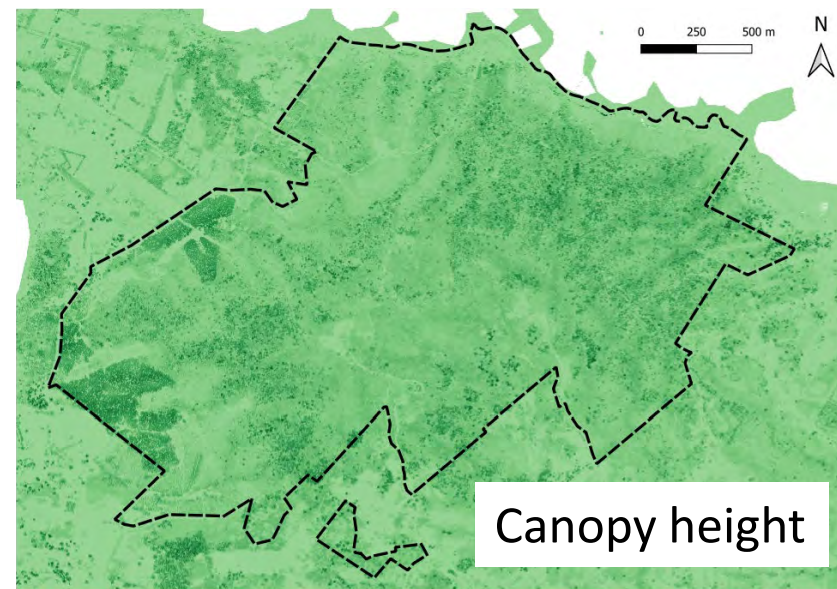


# Environmental variables



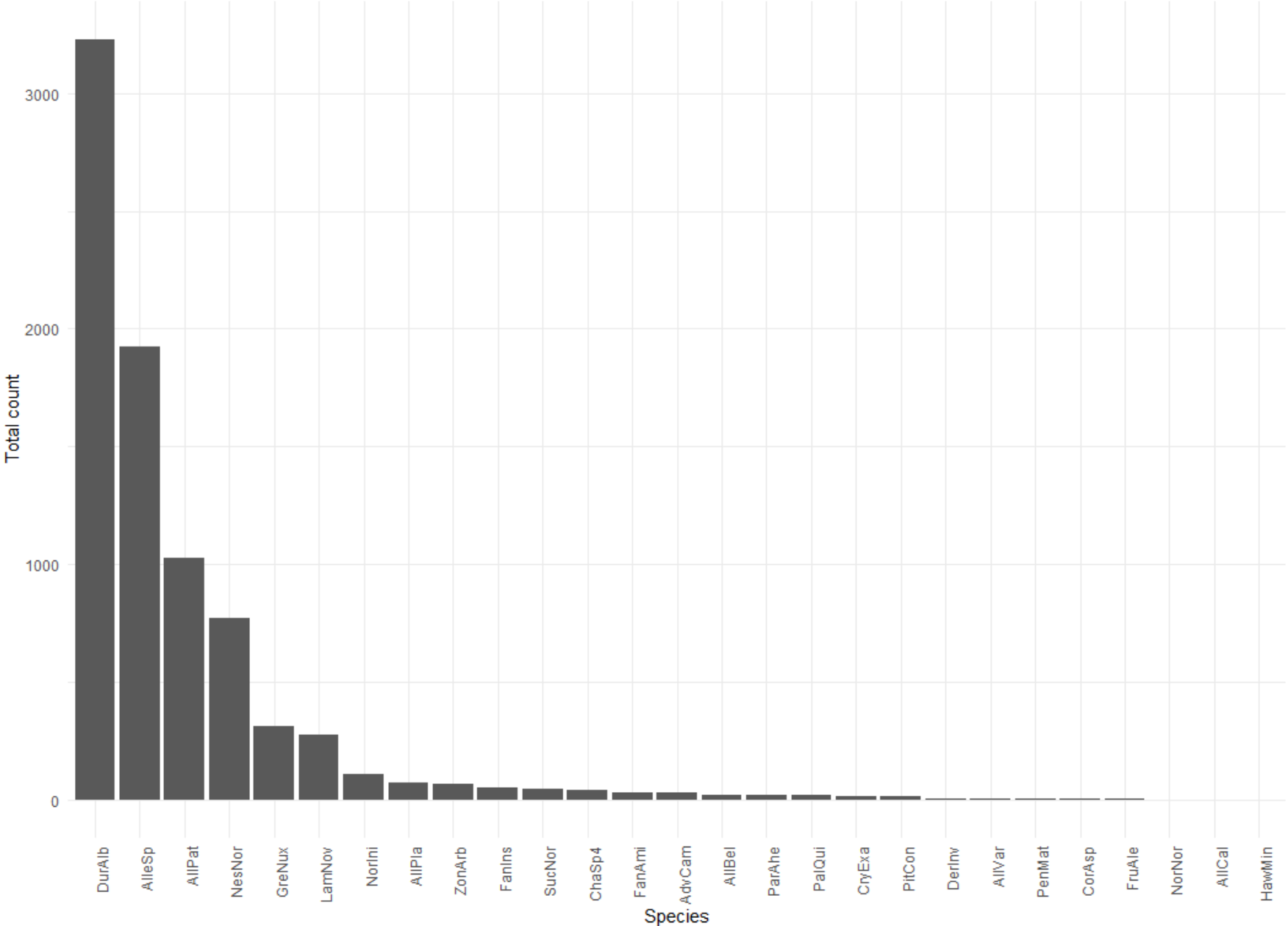


# Environmental variables





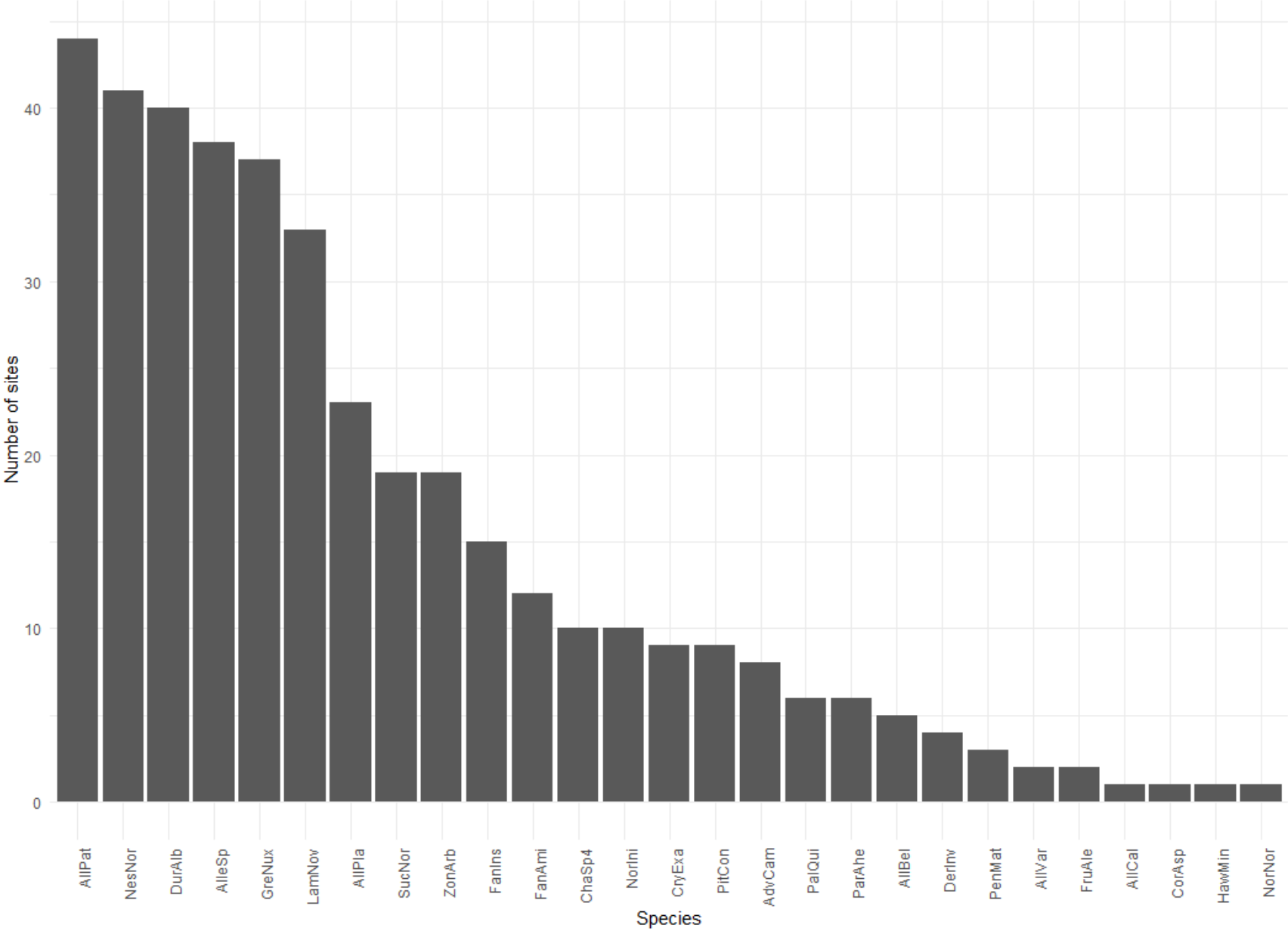
Abundance per species



Exploratory  
statistics



Number of sites per species



Exploratory  
statistics

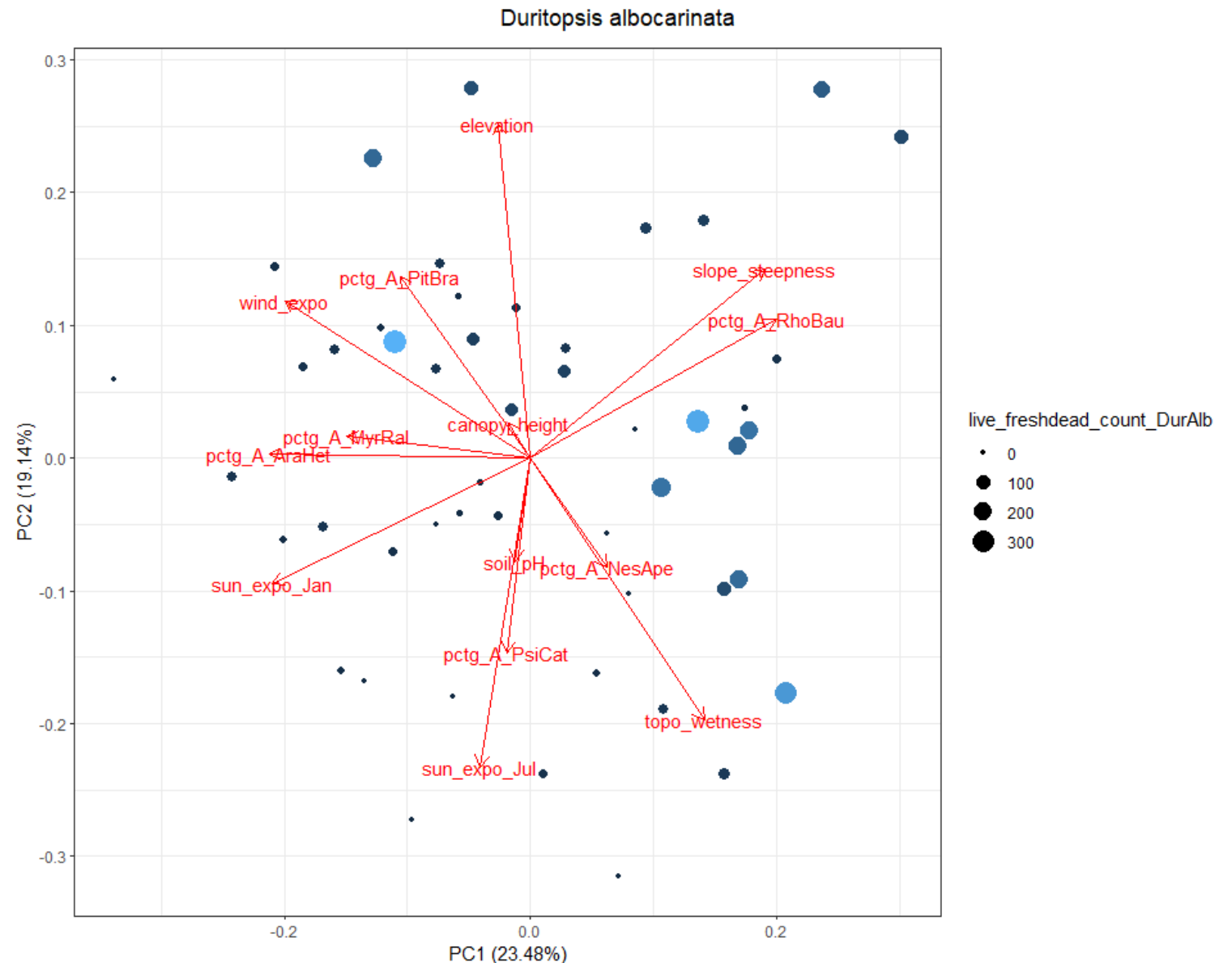


# Statistical analyses

Principal Component Analysis (PCA) – to identify variables that explain the most differences between sites



*Duritopsis albocarinata*  
CC-BY-SA Museum of New Zealand  
Te Papa Tongarewa



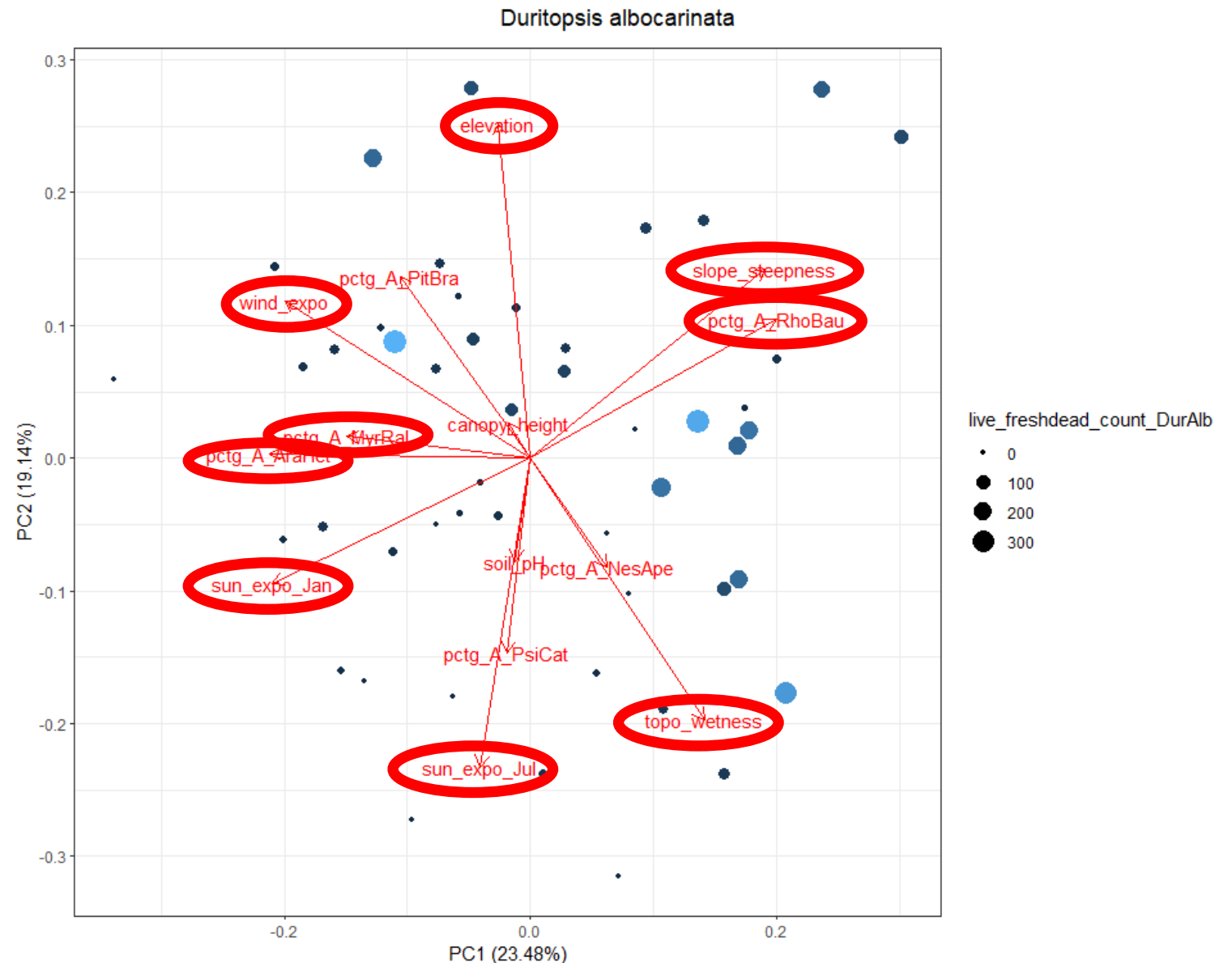


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# Statistical analyses

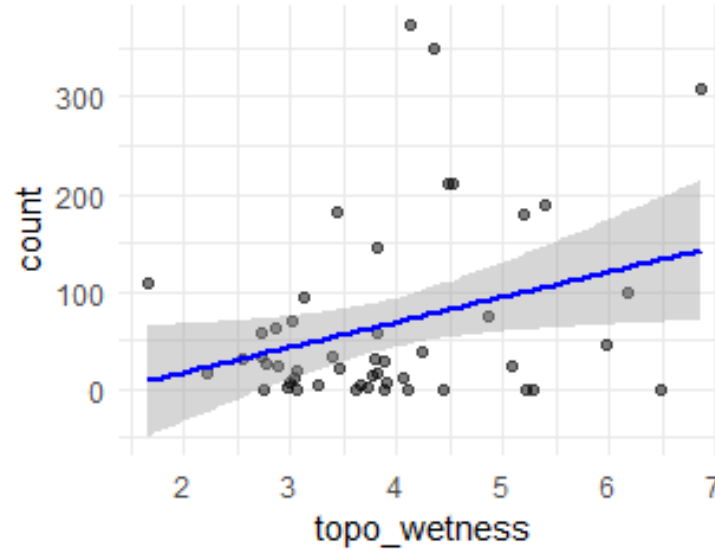
Generalised Additive Model  
(GAM) – to model species  
abundance against the key  
variables



# Statistical analyses

Generalised Additive Model (GAM) – to model species abundance against the key variables

Topo. wetness vs. snail abundance

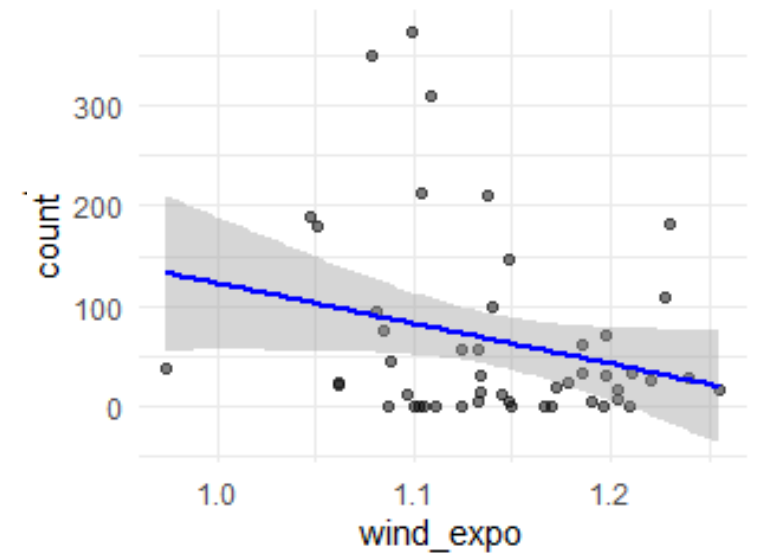
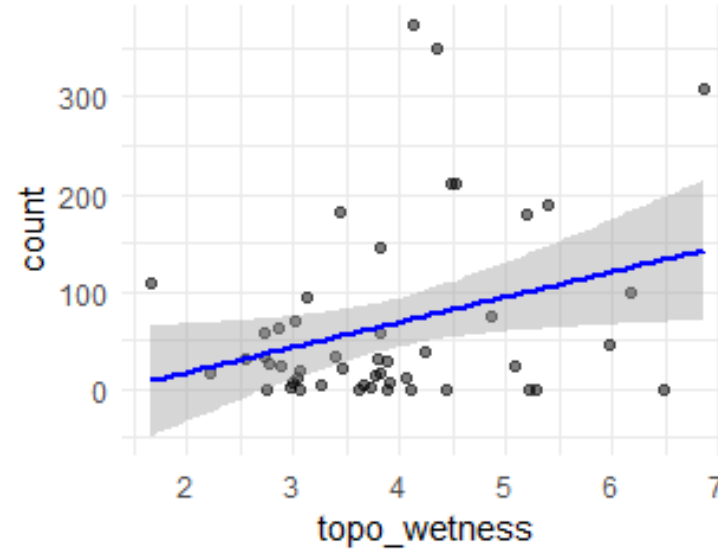




# Statistical analyses

Generalised Additive Model (GAM) – to model species abundance against the key variables

Topo. wetness vs. snail abundance    Wind exposure vs. snail abundance

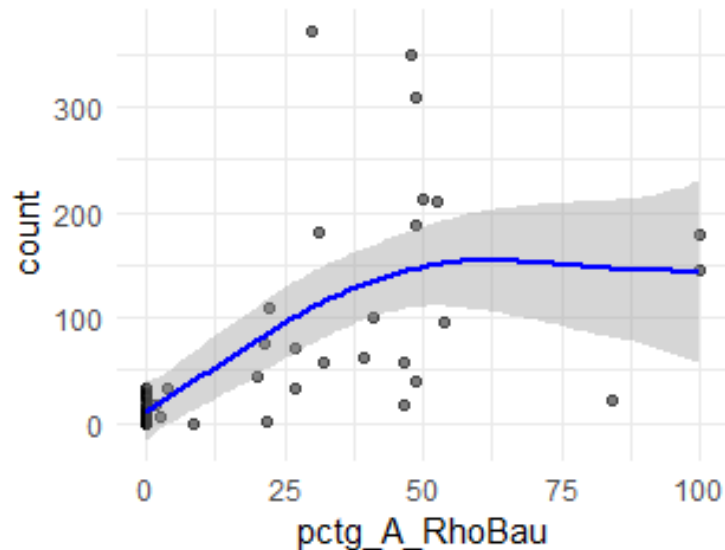




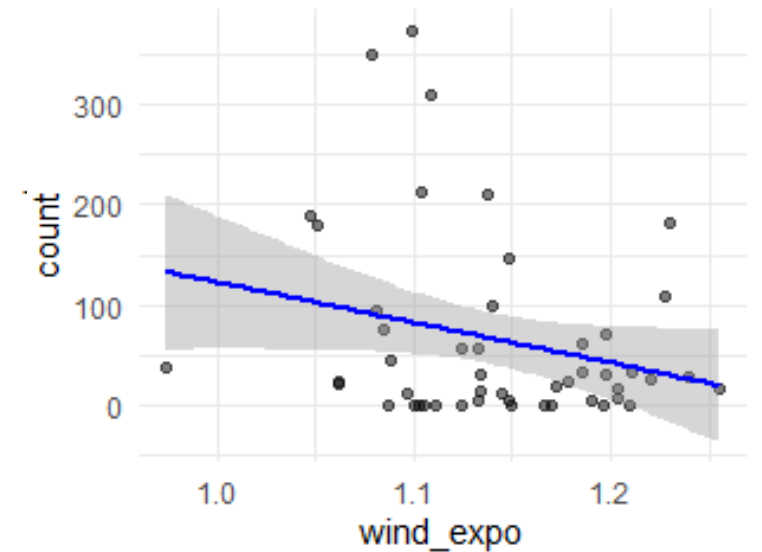
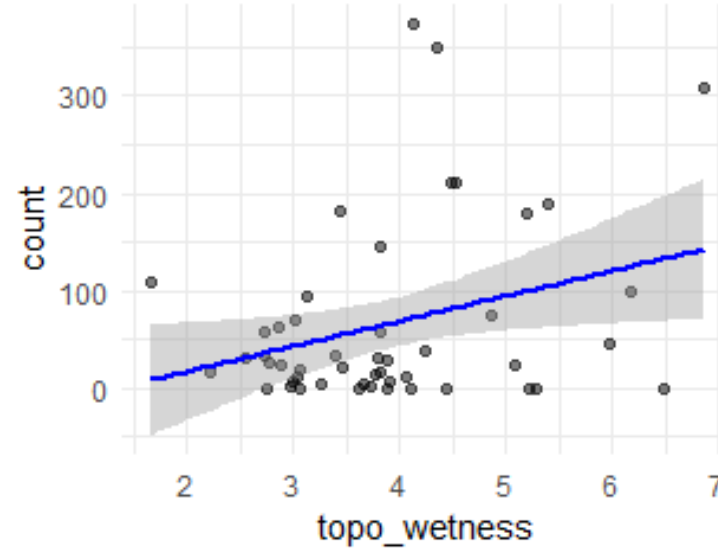
# Statistical analyses

Generalised Additive Model (GAM) – to model species abundance against the key variables

Palm % cover vs. snail abundance



Topo. wetness vs. snail abundance    Wind exposure vs. snail abundance

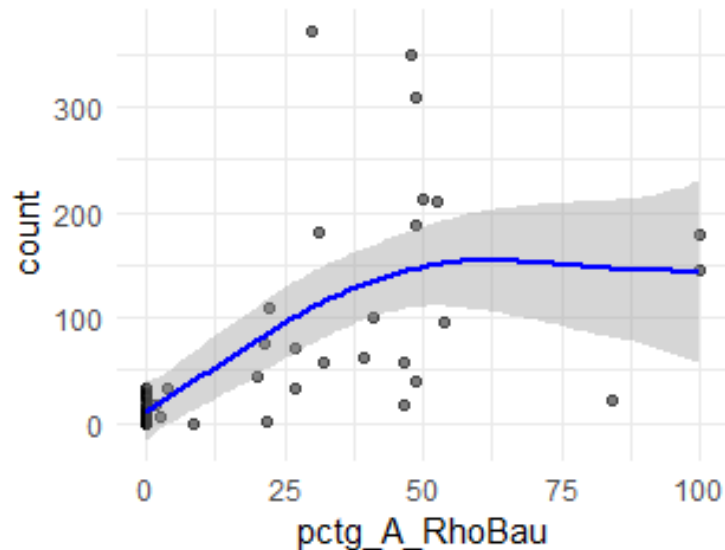




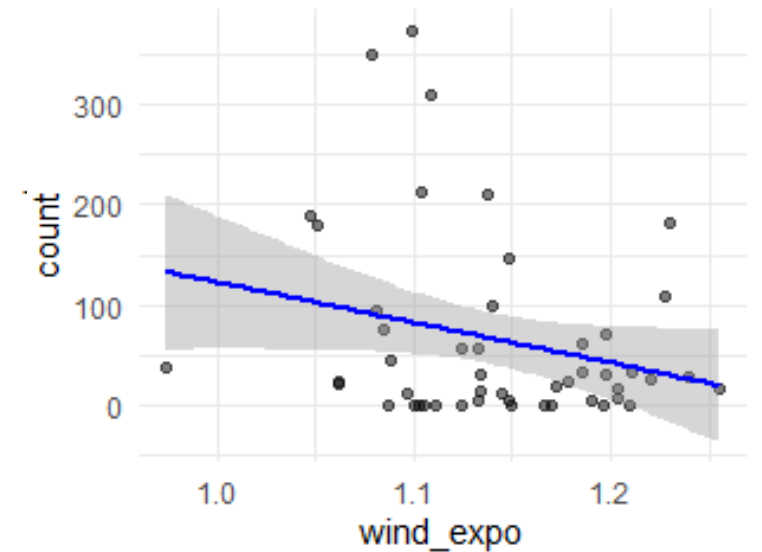
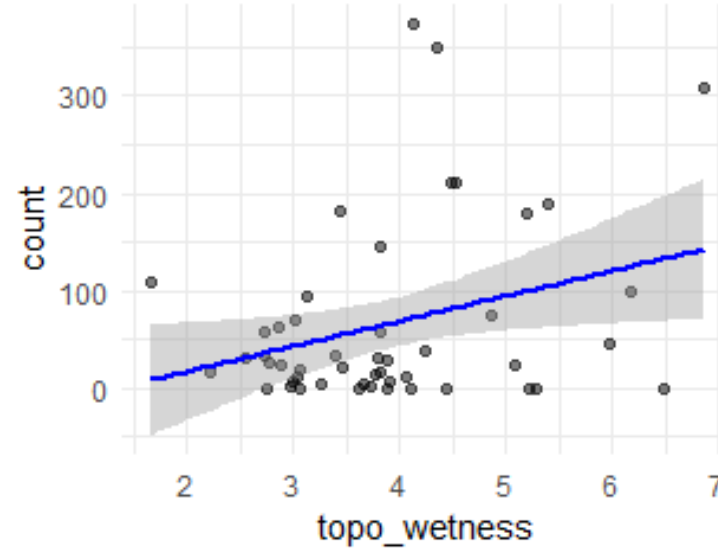
# Statistical analyses

Generalised Additive Model (GAM) – to model species abundance against the key variables

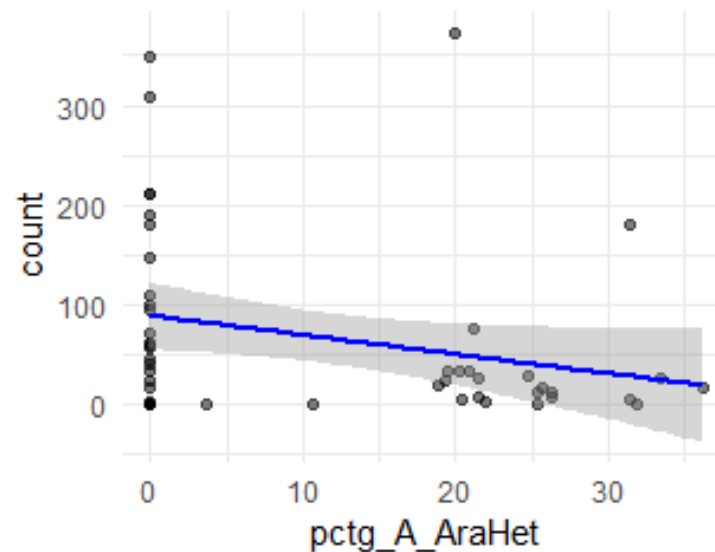
Palm % cover vs. snail abundance



Topo. wetness vs. snail abundance    Wind exposure vs. snail abundance



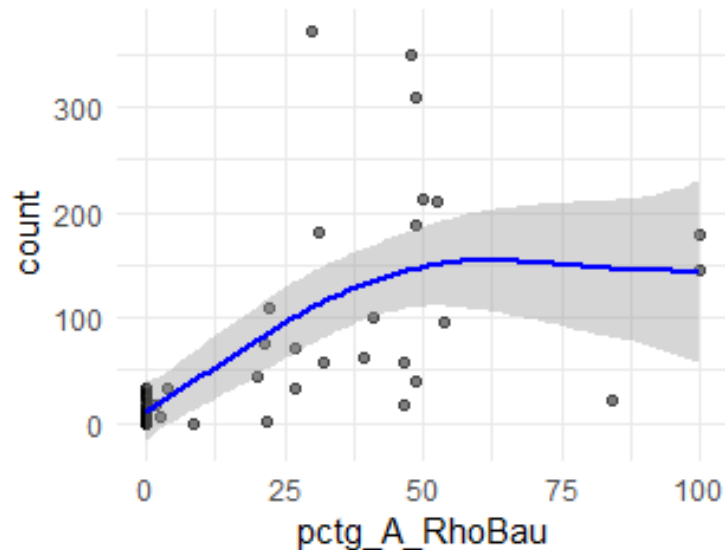
Pine % cover vs. snail abundance



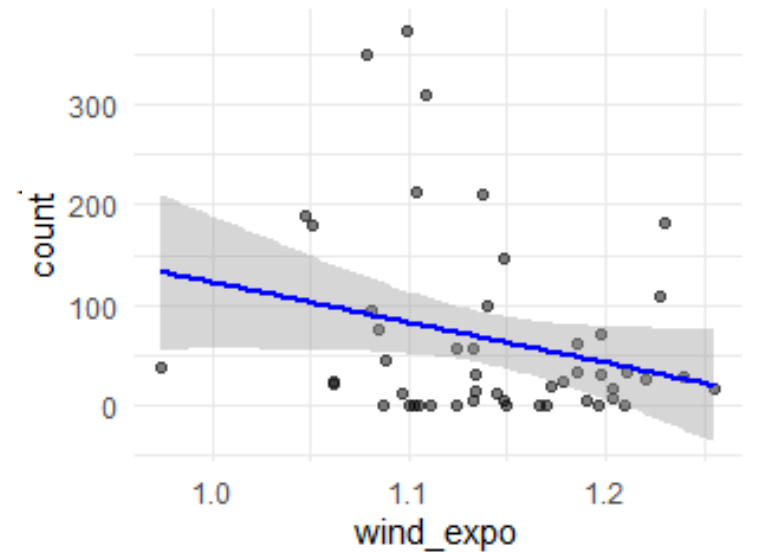
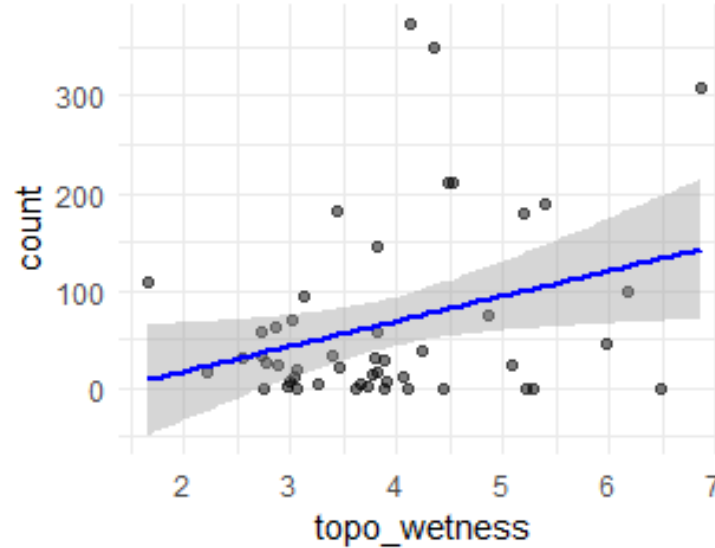
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Generalised Additive Model (GAM) – to model species abundance against the key variables

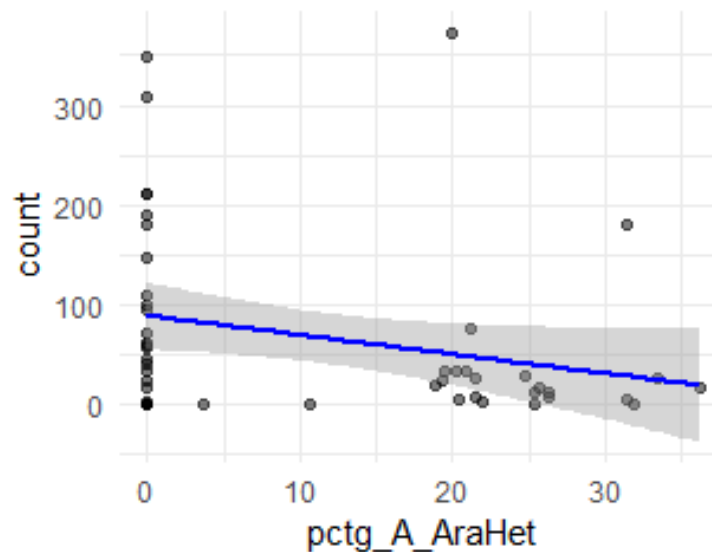
Palm % cover vs. snail abundance



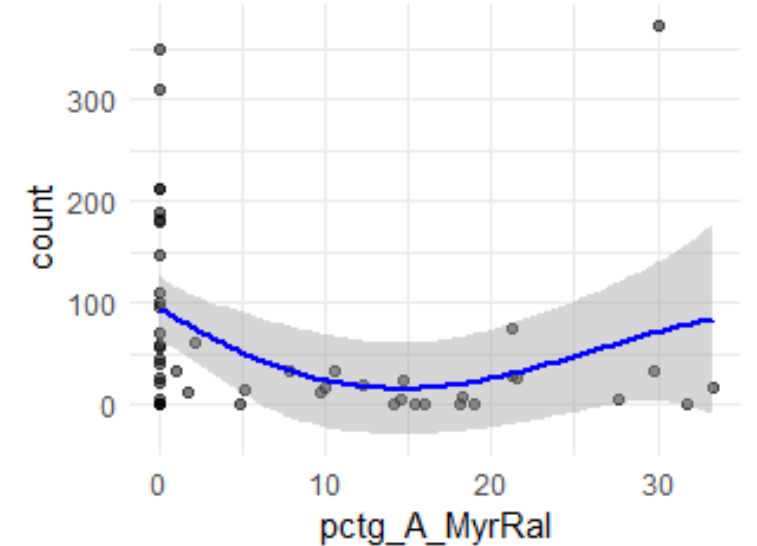
Topo. wetness vs. snail abundance      Wind exposure vs. snail abundance



Pine % cover vs. snail abundance

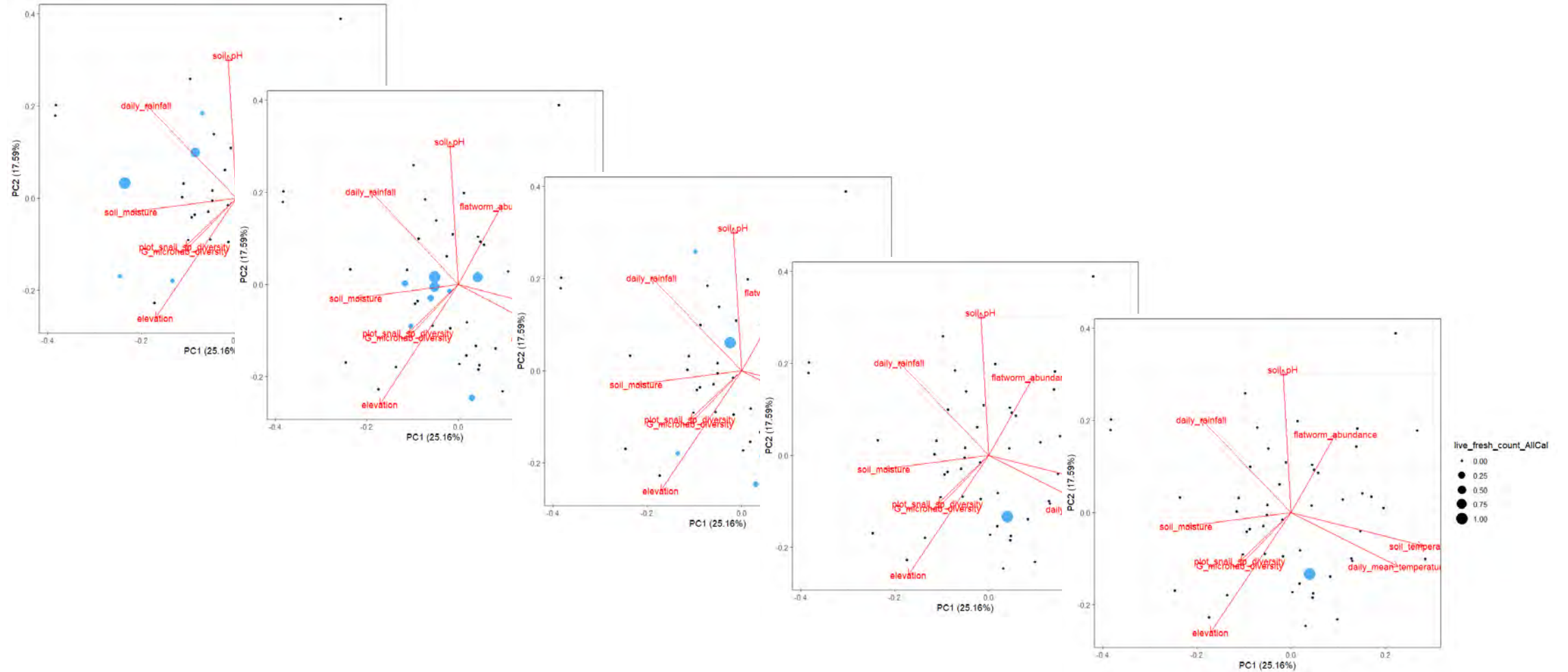


Beech % cover vs. snail abundance





# Repeat the analyses for all other land snail species



## Question 2

Where are the species distributed?



# Data quality check

Field survey data

Museum data

Verify records

- Species identities
- Time of collection
- Specimen condition
- Geographical coordinates

# Data quality check

Field survey data

Museum data

Verify records

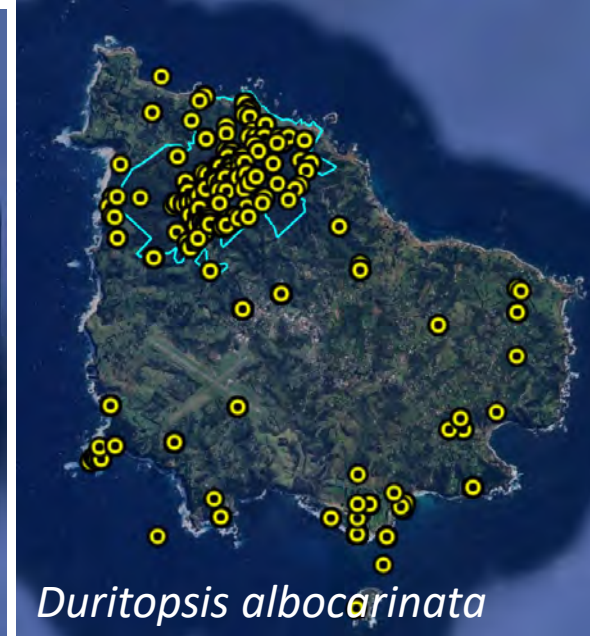
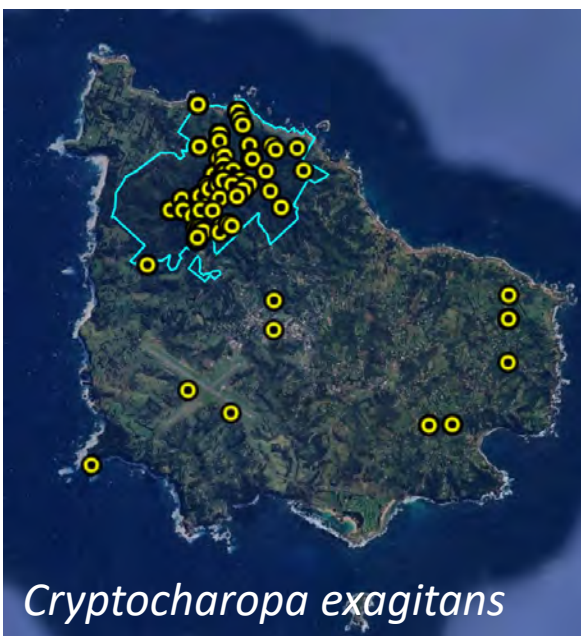
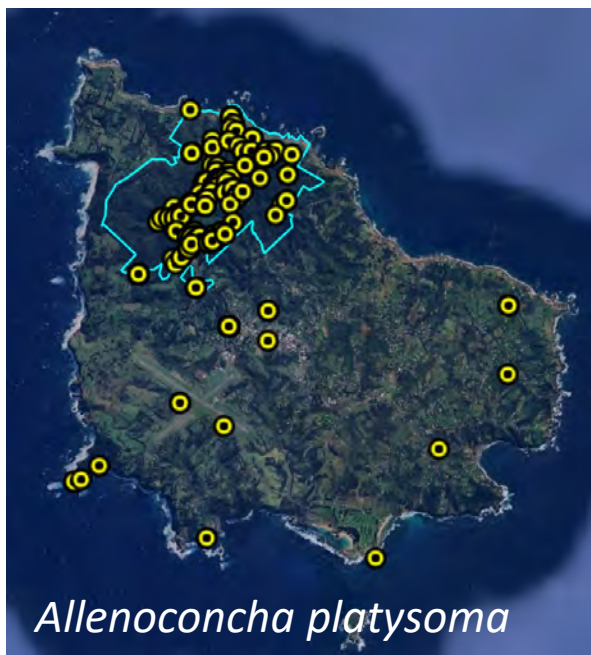
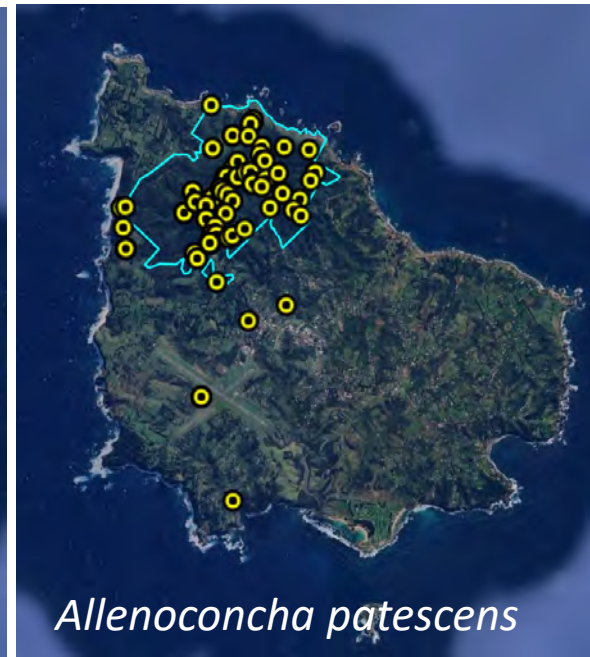
- Species identities
- Time of collection
- Specimen condition
- Geographical coordinates



Pearce (2008)

<http://dx.doi.org/10.4003/006.026.0211>





# Spatial analysis

Species Distribution Modelling



# Spatial analysis

## Species Distribution Modelling

Example – Mt Kaala, New Caledonia



*Arthropodium sp.nov.*



*Burretiokentia hapala*



*Casearia kaalaensis*



*Cleidion velutinum*



*Comptonella fruticosa*



*Deltaria brachyblastophora*



*Gynochthodes truncata*



*Myrsine novocaledonica*  
subsp. *kaalaensis*

# Spatial analysis

## Identify Important Land Snail Conservation Areas

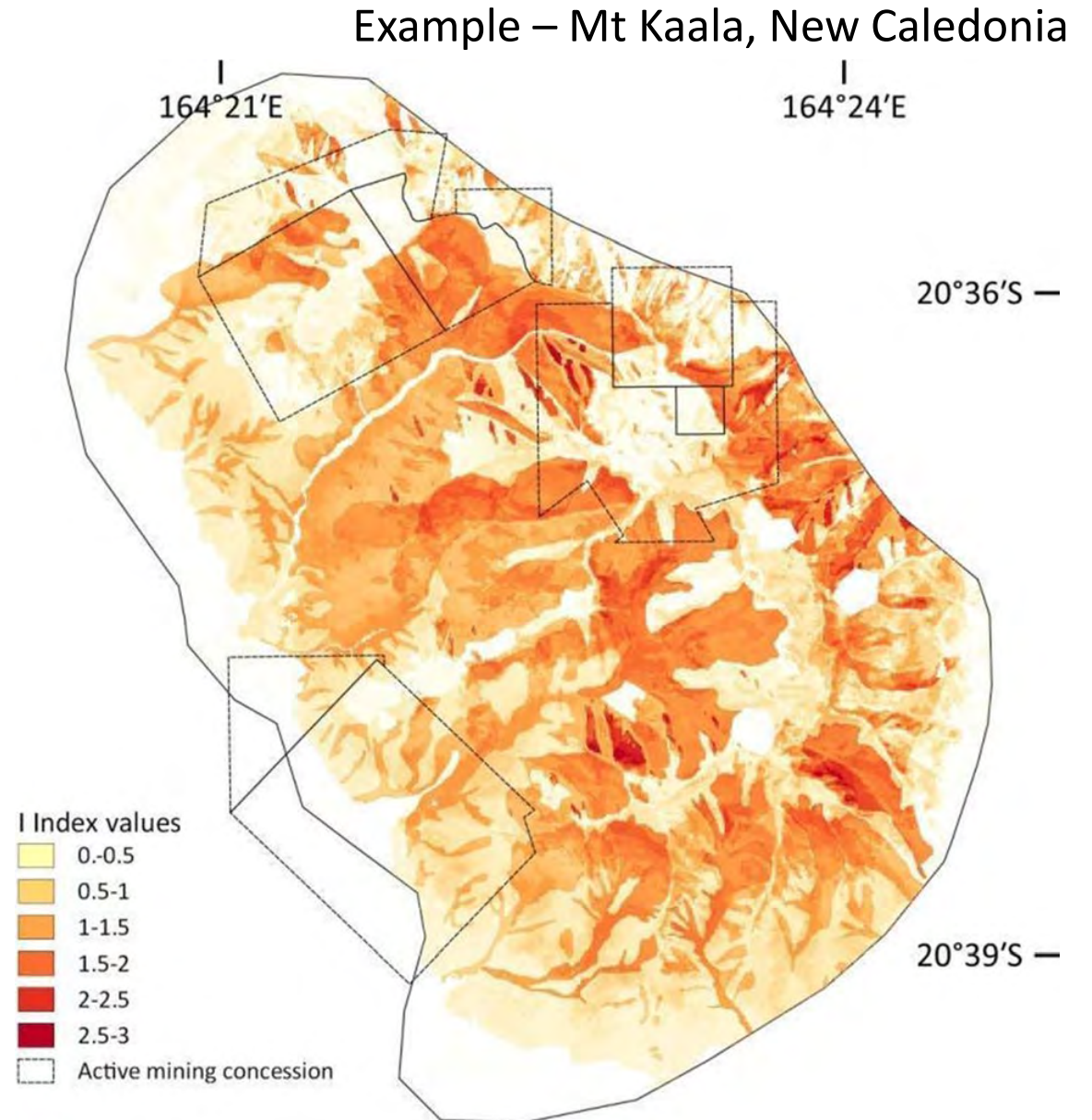
- Density of endemic species
- Conservation priority index score
- % overlap with protected areas  
and land uses



# Spatial analysis

## Identify Important Land Snail Conservation Areas

- Density of endemic species
- Conservation priority index score
- % overlap with protected areas and land uses





# Implications

- Prioritise conservation management in areas of important land snail habitats
- Restore degraded habitats for native snail reintroduction





# Acknowledgements

Hoong Fatt Foon | Melinda Wilson | Nigel Greenup | Sara Freeland | Lilli King | Tara Patel  
Mark Scott | Mark Hallam | Rob Varman | The Norfolk Island Community  
Norfolk Island Flora & Fauna Society

We acknowledge and pay our respects to the Norfolk Islanders, particularly to the Norf'k Ailen Kaunsl' Eldas and all Pitkern descendants who call Norf'k home, as the traditional owners of Norfolk Island where we conducted this research.

We acknowledge the Dharug, Dharawal, Gandangarra, Eora and Wiradjuri Peoples as the First Peoples and Traditional Custodians of the land and waterways on which Western Sydney University and Australian Museum stands, and pay our respects to their elders past, present, and future.



Australian Government  
Parks Australia



Australian Government  
Department of Climate Change, Energy,  
the Environment and Water

UNITAS  
MALACOLOGICA



WESTERN SYDNEY  
UNIVERSITY





*Thank you!*

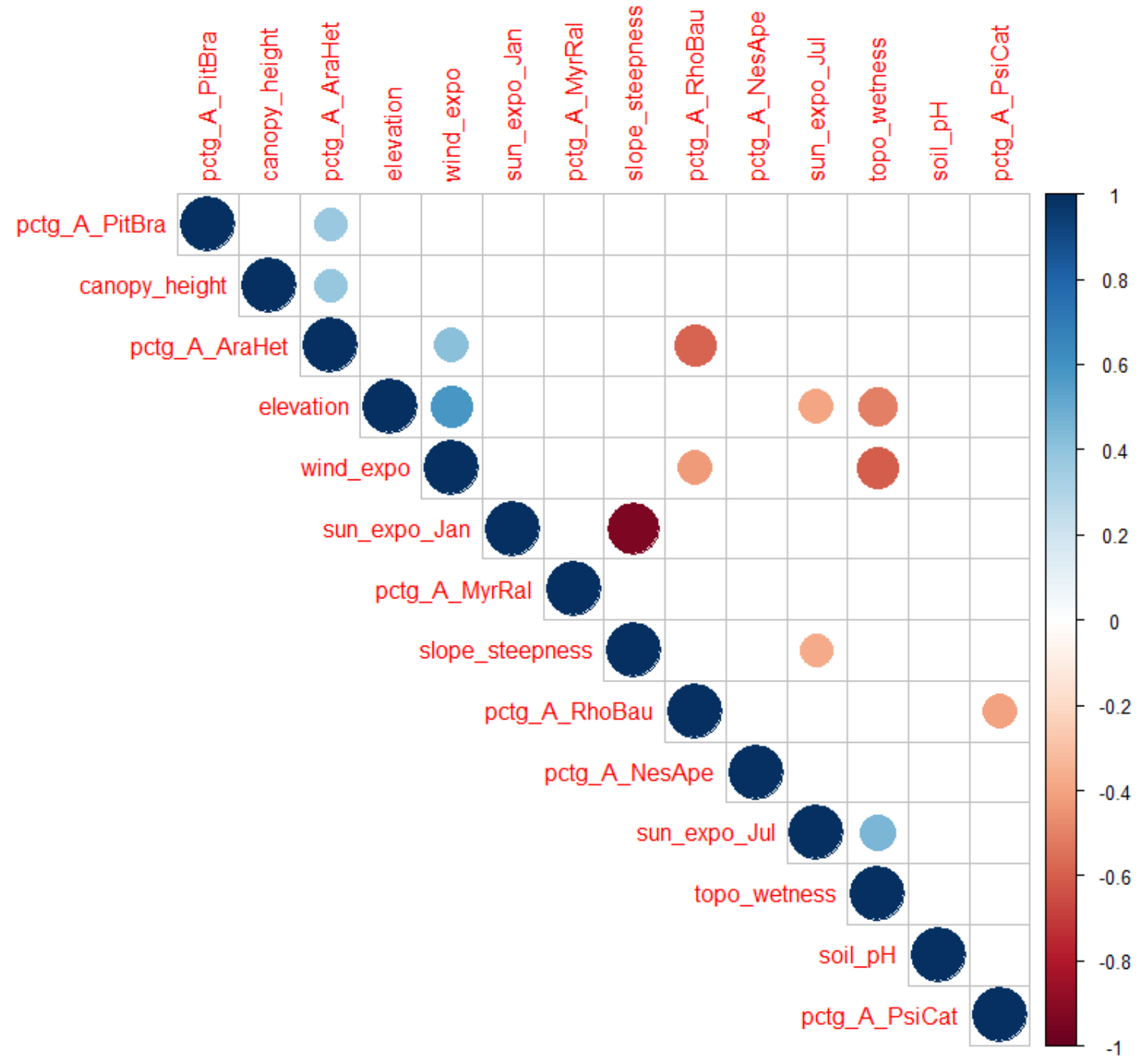




Appendix slides

# Statistical analyses

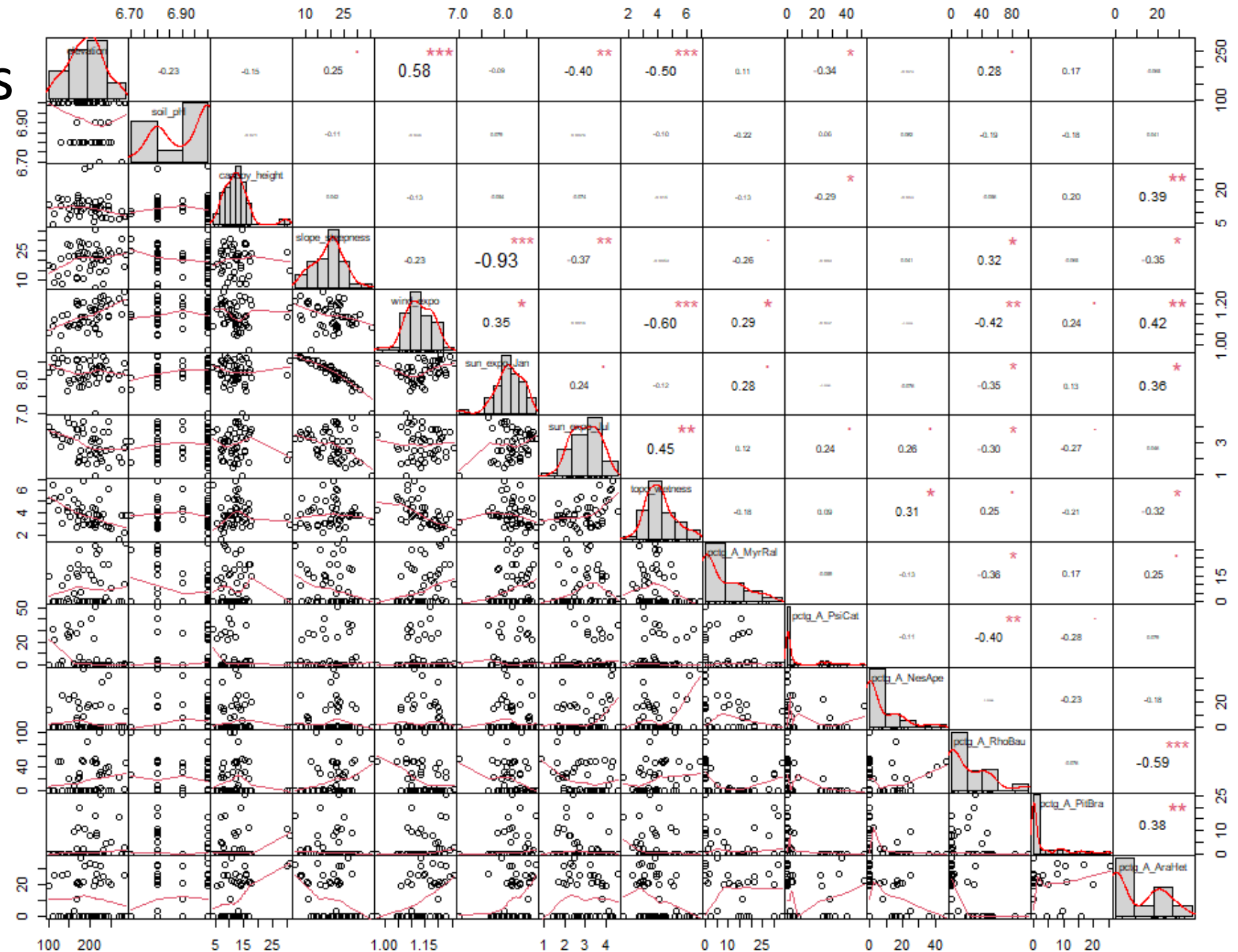
Exploratory – correlation plots





# Statistical analyses

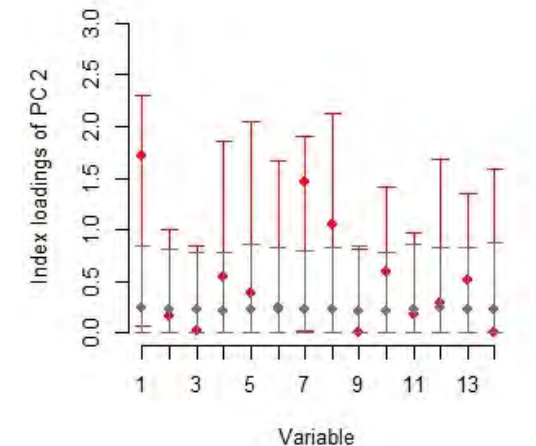
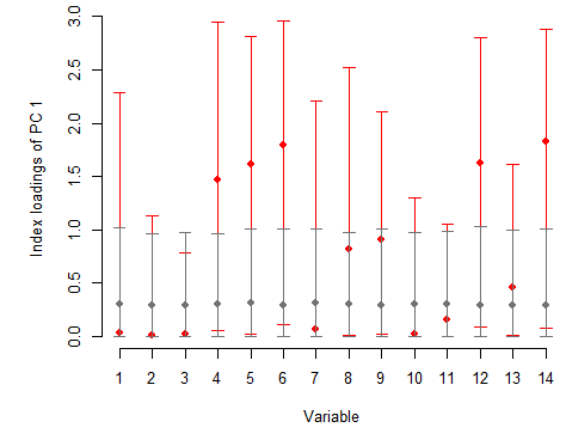
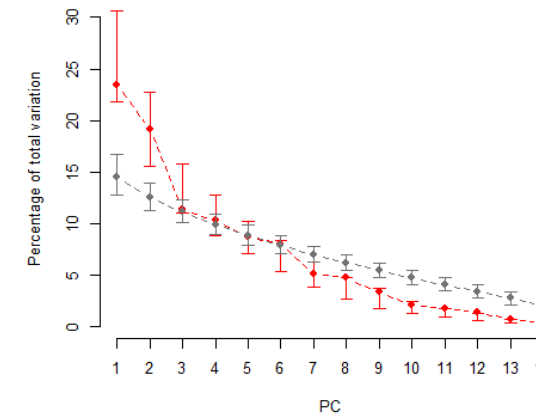
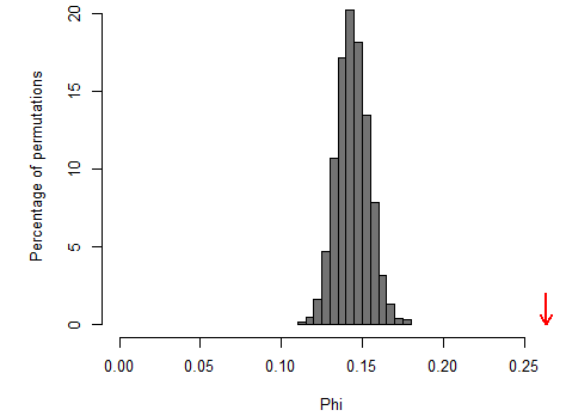
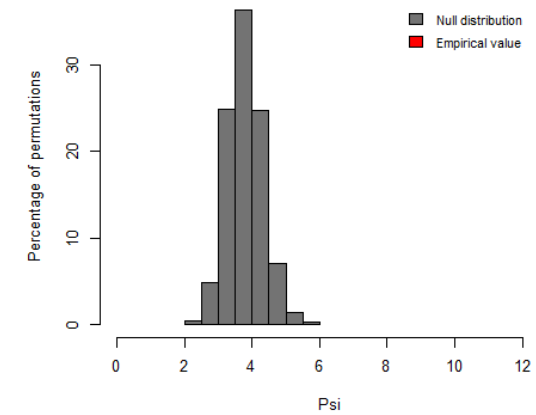
Exploratory – correlation plots



# Statistical analyses

PCA – to determine variables that explain the main differences between sites

	PC1	PC2
Eigenvalues	3.28650697	2.67975565
Percentage variance	23.4750498	19.1411118
Cumulative variation	23.4750498	42.6161616
Macrohabitat variables	Correlation (rS)	Correlation (rS)
elevation	0.09136752	-0.80157926*
soil pH	0.04578524	0.24703297
canopy height (m)	0.06443903	-0.08498624
slope steepness	-0.66935947*	-0.4520428
wind exposure	0.70172922*	-0.37691269
sun exposure in January (kWh m2)	0.73847128*	0.30237588
sun exposure in July (kWh m2)	0.14417835	0.74013354*
topographic wetness index	-0.49660362	0.62776845*
percentage cover Myrsine ralstoniae	0.52592833*	-0.052473
percentage cover Psidium cattleianum	0.06863041	0.4682351
percentage cover Nestegis apetalum	-0.21702941	0.26126694
percentage cover Rhopalostylis baueri	-0.7027268*	-0.33169891
percentage cover Pittosporum bracteolatum	0.37320006	-0.43458307
percentage cover Araucaria heterophylla	0.74644903*	-0.01029147





# Statistical analyses

GAM – to model species abundance against the key variables

Family: poisson  
Link function: log

Formula:

```
live_freshdead_count_DurAlb ~ s(elevation, k = 10) + s(slope_steepness,
  k = 10) + s(wind_expo, k = 10) + s(topo_wetness, k = 10) +
  s(sun_expo_Jan, k = 10) + s(sun_expo_Jul, k = 10) + s(pctg_A_MyrRal,
  k = 10) + s(pctg_A_RhoBau, k = 10) + s(pctg_A_AraHet, k = 10)
```

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	2.7372	0.1169	23.41	<2e-16 ***

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Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

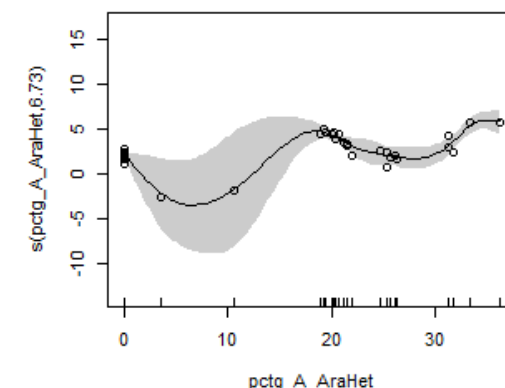
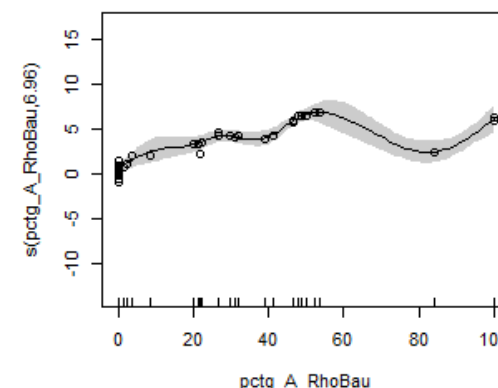
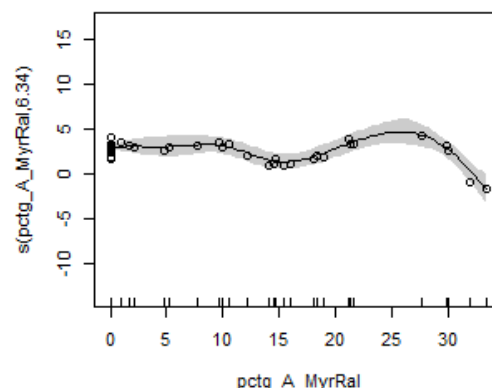
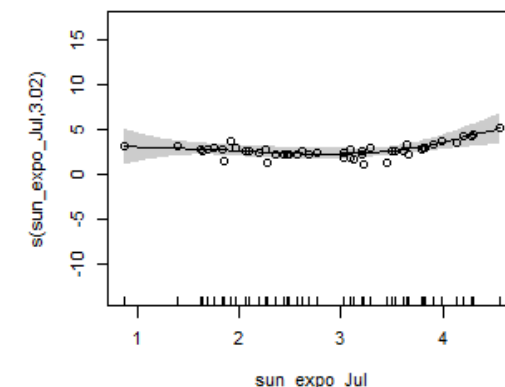
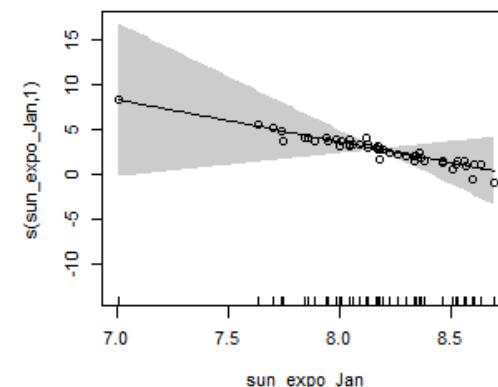
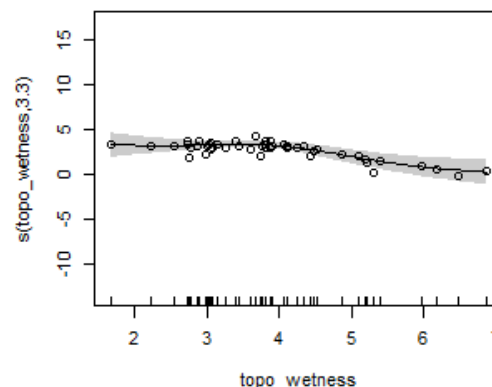
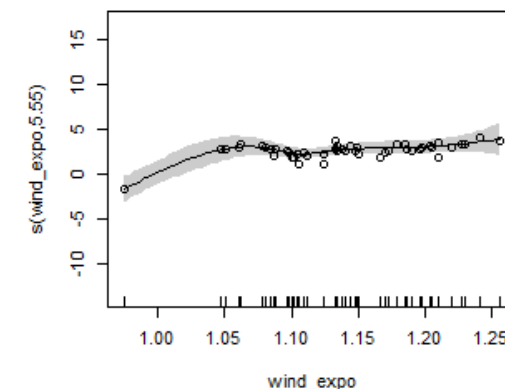
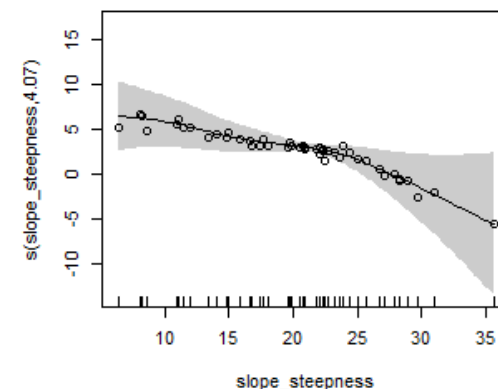
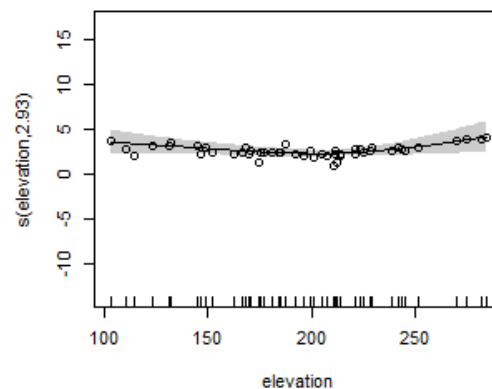
	edf	Ref.df	Chi.sq	p-value
s(elevation)	2.933	3.147	1.344	0.7251
s(slope_steepness)	4.069	4.352	6.707	0.2205
s(wind_expo)	5.553	5.976	30.952	3.21e-05 ***
s(topo_wetness)	3.298	3.524	10.877	0.0101 *
s(sun_expo_Jan)	1.000	1.000	1.652	0.1987
s(sun_expo_Jul)	3.017	3.261	6.940	0.1216
s(pctg_A_MyrRal)	6.344	6.744	48.180	< 2e-16 ***
s(pctg_A_RhoBau)	6.959	7.219	196.059	< 2e-16 ***
s(pctg_A_AraHet)	6.733	7.307	71.051	< 2e-16 ***

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Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.999 Deviance explained = 99.7%

-REML = 233.19 Scale est. = 1 n = 49



# Statistical analyses

GAM – to model species abundance against the key variables

*Duritopsis albocarinata* counts versus macro-habitat variables

