Biodiversity across Scales: Mapping taxonomic, phylogenetic, and functional diversity with eDNA, field surveys, and remote sensing



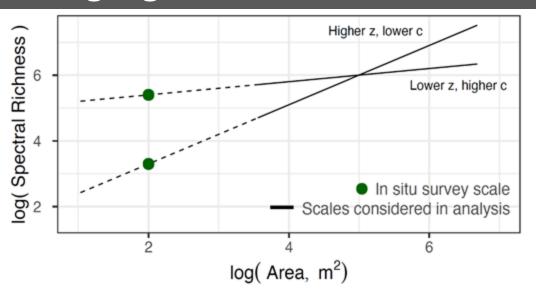
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Objectives

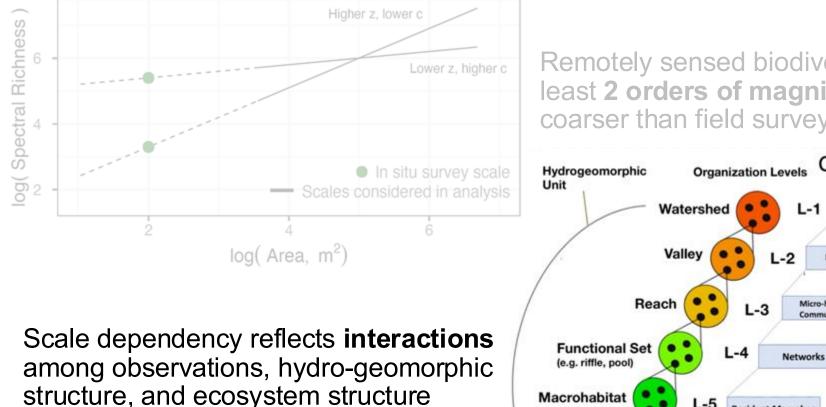
Question	Significance	
RELATION: How are phylogenetic, taxonomic, and functional diversity related to each other?	Map biodiversity consistently, globally, and at regular intervals using remote sensing	
SPACE: How does the hydrologic structure of watersheds organize phylogenetic, taxonomic, and functional diversity spatially?	As watersheds change and/or are modified, spatial patterns in biodiversity are also expected to change.	
TIME: How do hydroclimatic processes influence the temporal signal of phylogenetic, taxonomic, and functional diversity?	Constrain when and how frequently we need measurements of biodiversity using eDNA and remote sensing	

Bridging observations across scales



Remotely sensed biodiversity is at least **2 orders of magnitude** coarser than field surveys

Bridging observations across scales



Microhabitat

Remotely sensed biodiversity is at least 2 orders of magnitude coarser than field surveys

L-1

Macrosystem

Measured

Measured with eDNA

with Remote Sensing

Landscape

Micro-habitat

Communities

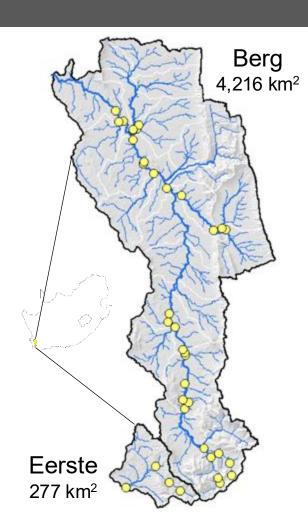
Resident Macrobes

Microbial Guilds

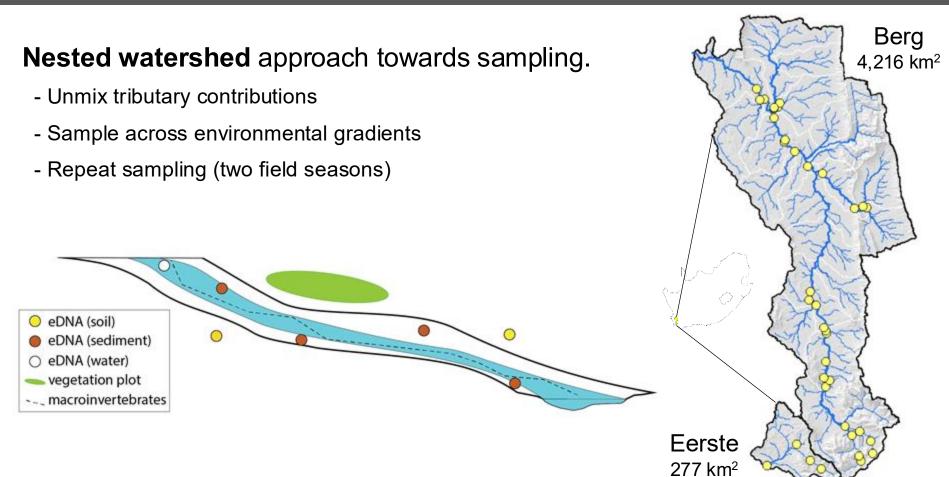
Approach

Nested watershed approach towards sampling.

- Unmix tributary contributions
- Sample across environmental gradients
- Repeat sampling (two field seasons)

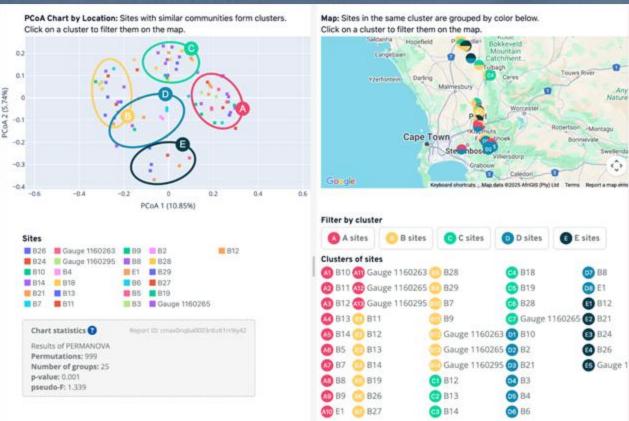


Approach



environmental DNA

eDNA Explorer



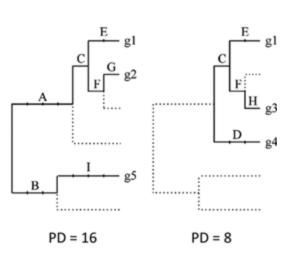
Current status:

- 560 samples uploaded into eDNA Explorer
- Working with end-users on permissions and usability of interface
- Distinct fungal, eukaryotic, and bacterial communities as function of distance from headwaters and human modification index

environmental DNA

Ongoing:

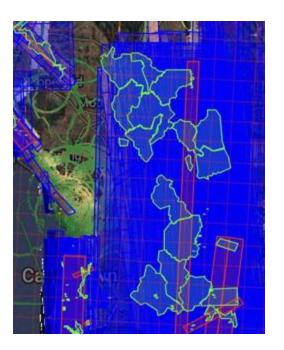
1. Analyze **phylogenetic diversity** within eDNA Explorer



2. Incorporate other variables like **invasive species cover** and **stream morphology**



3. Integrate with **functional diversity** metrics



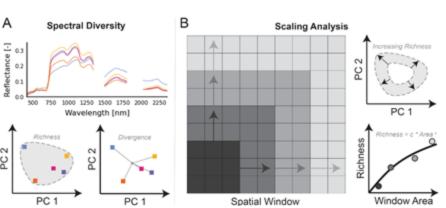
Spectral and functional diversity



Spectral PCA

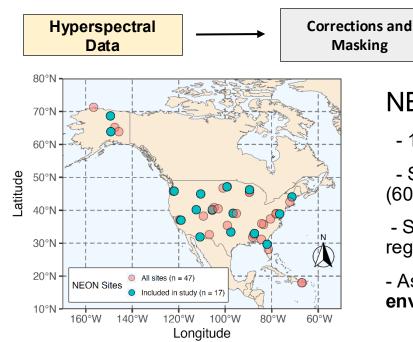
Current status:

- Built cloud-based computational workflow using v1 BRDF/Topo corrected reflectance (aviris-ng) and foliar traits
- Berge/Eerste analysis will use v2 data products
- Approach: Calculate spectral and functional richness at different spatial scales to characterize scaling relationship



Spectral and functional diversity

	Wavelengths	Spectral resolution	Spatial resolution	Temporal resolution
AVIRIS-NG	380 - 2510 nm	5 nm	5-7 m approx.	Single pass
NEON AOP	380 - 2500 nm	10 nm	1-3 m approx.	Single pass



NEON Analysis

- 15 sites (105 plots) representing large range of ecosystems

Dimensionality

Reduction

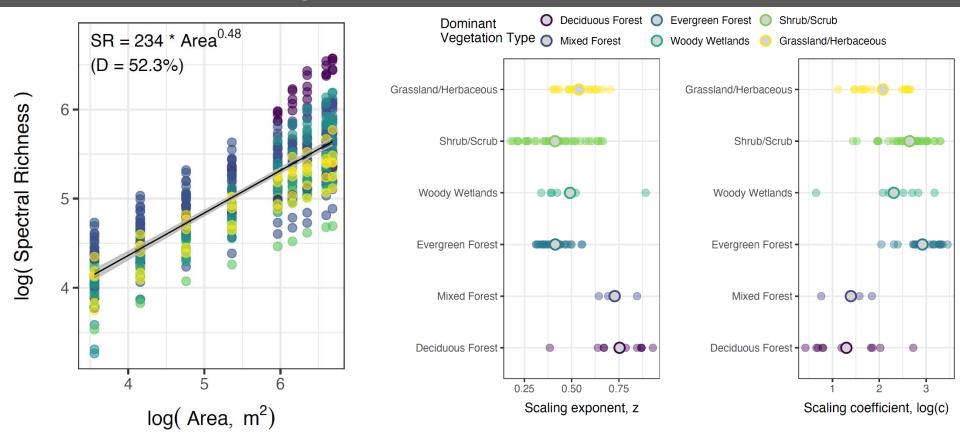
Compute

Functional

Metrics

- Spectral richness calculated over large range of window sizes (60 to 2,000 m wide kernel)
- Scaling relationships characterized using **best-fit power law** regression
- Assess how regression parameters relate to other potential **environmental covariates** (e.g., climate, topography, landuse)

Spectral diversity of NEON sites



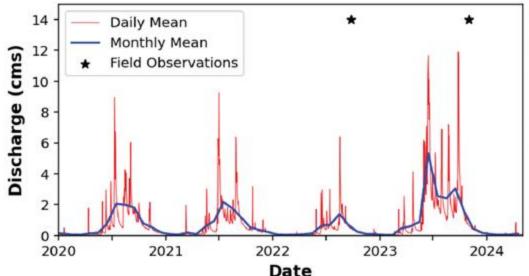
Nonlinear scaling robust but intra-site variations on par with inter-site ones

Opportunities in the Berg/Eerste systems

Historic floods (and debris flows) along Berg in between field seasons

 Test how diversity signals are to temporal disturbances







Opportunities in the Berg/Eerste systems

Functional Process Zones: river reaches with similar hydrological, geomorphological, and ecological traits

Test how robust diversity signals are to spatial patterning set up by the river network

Derived using OpenRES, a QGIS plugin our team is developing to classify FPZs:

OpenRES extracts 9 key variables for each user-defined stream segment:

ELE: Elevation

PRE: Mean Annual Precipitation

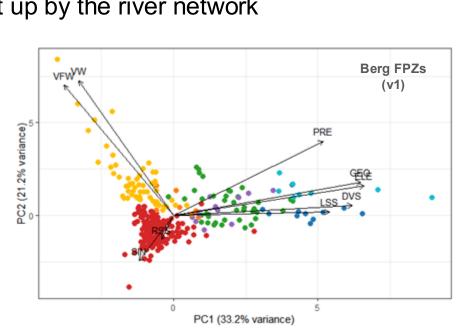
GEO: Geology

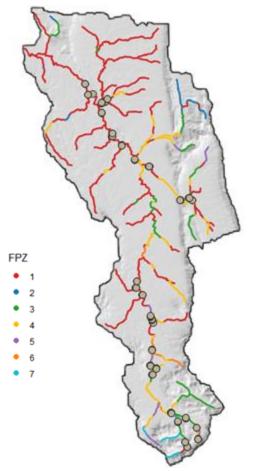
VFW: Valley floor width VW: Valley width LSS: Left side slope

RSS: Right side slope DVS: Down valley slope

SIN: Sinuosity

FPZs derived from unsupervised classification of these key variables





Much thanks to our sponsors and collaborators

