



easynem: an R package for computing and visualizing soil nematode ecological indices

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Introduction



Soil nematode communities

Data Preparation

Abundance Taxonomy Test design

↓ ↓ ↓

Genus	Family	Feeding	Cp_value
Aba...	Mer...	Herbivo...	1
Abb...	Phy...	Funivor...	3
Abe...	Enc...	Bacteri...	3

Data reading
read_nem()
read_nem2()

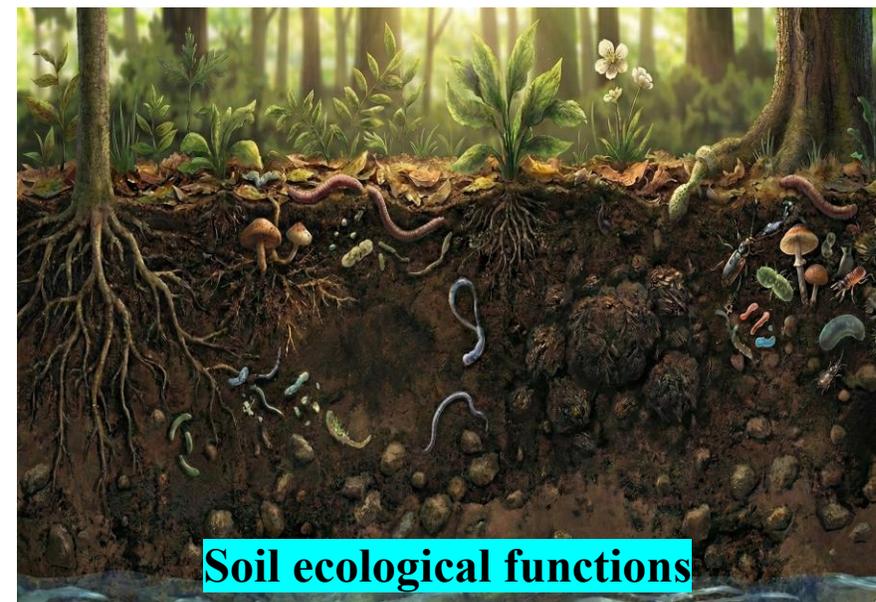
Automated linking to soil nematode databases

Database visualization
nem_database()

Data Pre-processing

Data filtering
filter_name()
filter_num()

Data transformation
trans_name()
trans_formula()
trans_norm()
trans_rare()
trans_combine()



Soil ecological functions

Data Visualization

nem_plot()

Life-history strategy Metabolic footprint Energy flows

Calculation of Nematode-based Indices

Nematode community

calc_alpha()
calc_beta()
calc_nemindex()
calc_ter()
calc_compare()

Nematode-based indices

calc_funguild()
calc_mf()
calc_ef()
calc_lm()



Highlights

```
# Installing and loading easynem
```

```
install.packages("easynem")
```

```
library(easynem)
```

```
# Import data
```

```
nem <- read_nem2(tab = nemtab,
```

```
  tax = nemtax,
```

```
  meta = nemmeta)
```

```
# Metabolic footprint analysis
```

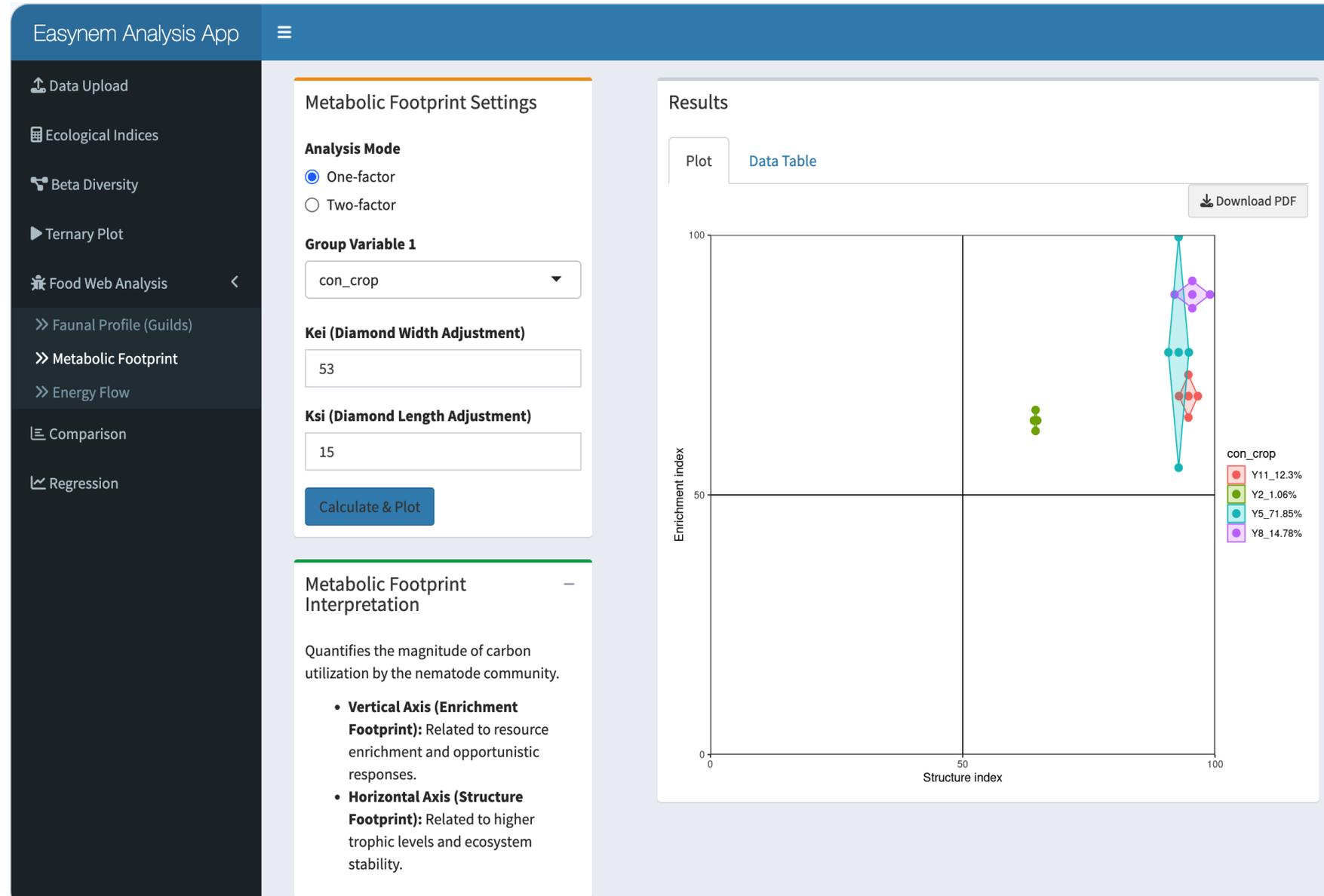
```
p <- nem |>
```

```
  calc_nemindex() |>
```

```
  calc_mf(con_crop) |>
```

```
  nem_plot(kei = 53, ksi = 15)
```

R Code



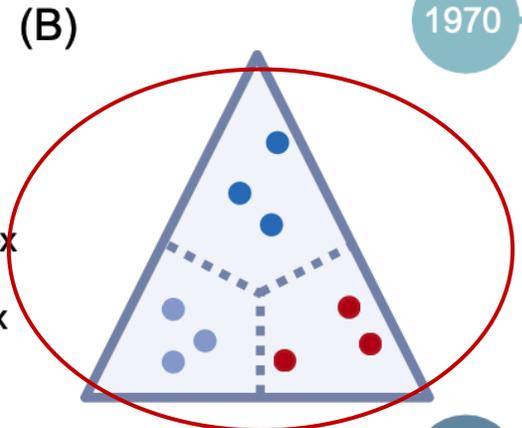
https://whkygl.shinyapps.io/Easynem_Analysis/



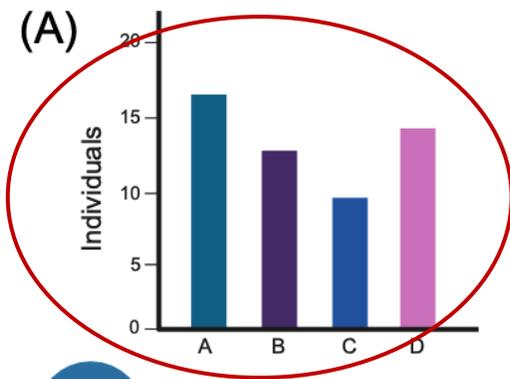
Overview of Nematode Ecological Indices

Life history strategies

- Cp-triangle
- Maturity Index 2-5
- Sigma Maturity Index
- Plant-Parasitic Index
- PPI/MI



1970



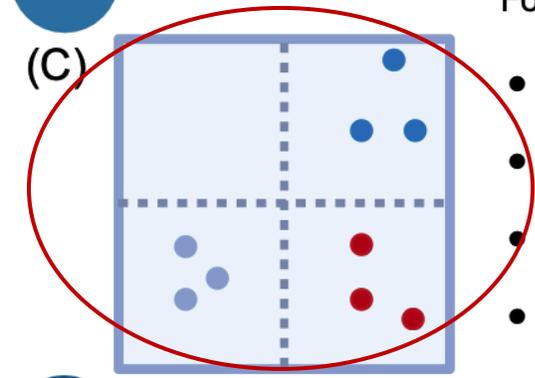
Abundance and diversity

- Species Richness (S)
- Simpson Index (D)
- Shannon Diversity Index (H')
- Pielou Evenness Index (J')

1990

Food web and faunal analysis

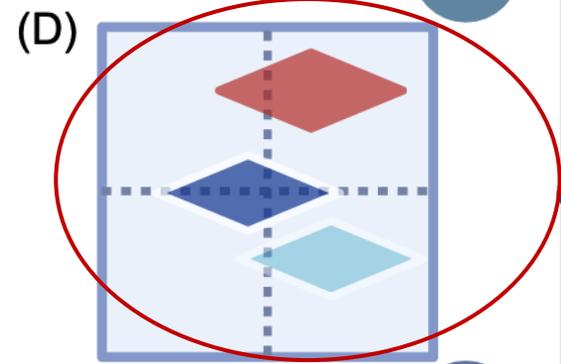
- Enrichment Index (EI)
- Structure Index (SI)
- Channel Index (CI)
- Basal Index (BI)



2000

Metabolic footprints

- Enrichment Footprint
- Structure Footprint
- Functional Footprint
- Aggregate Footprint
- Herbivore, Bacterial and Fungal Footprints

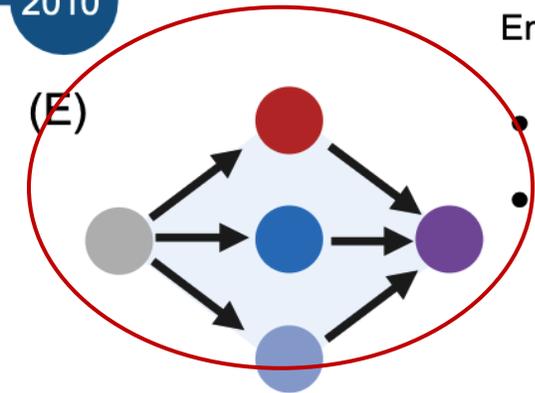


2010

Energetic structure

- Energy Flux
- Uniformity (U) of Soil
- Nematode Energetic Structure

NOW





Case 1: Trophic Groups and Life History Strategies



Kiwi orchard mulching experiment

CK: No crop cover

C2: 2 types of crop cover

C4: 4 types of crop cover

C8: 8 types of crop cover

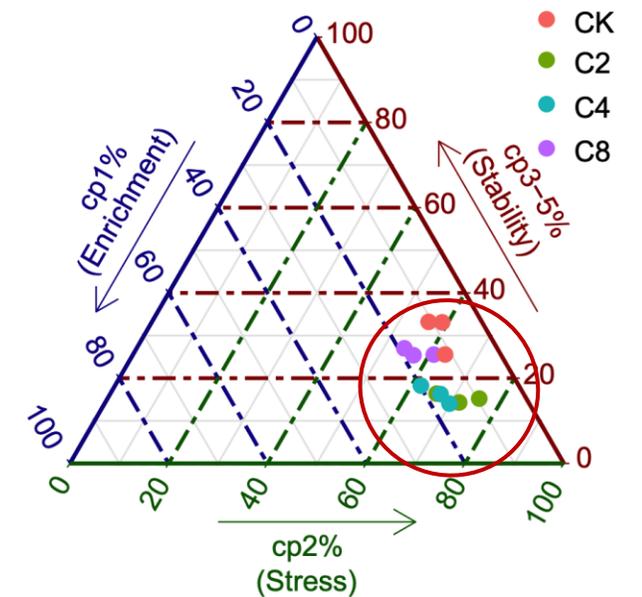
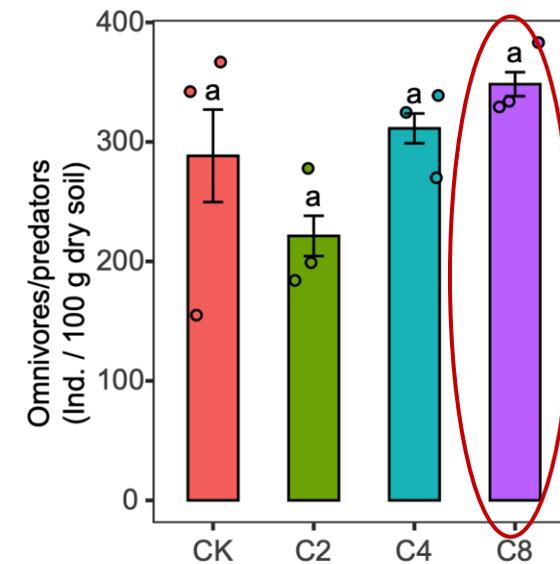
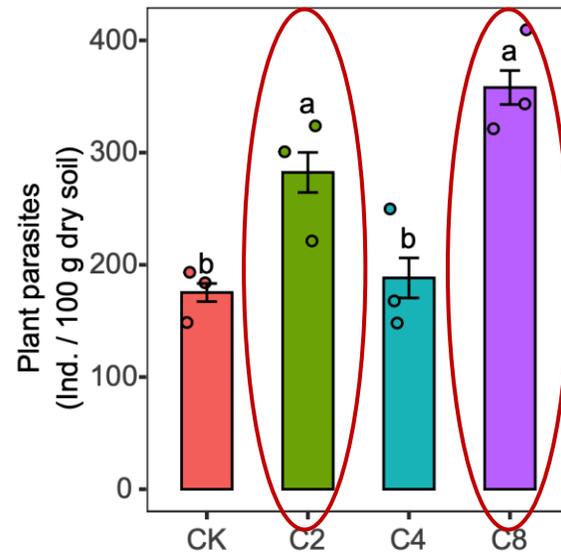
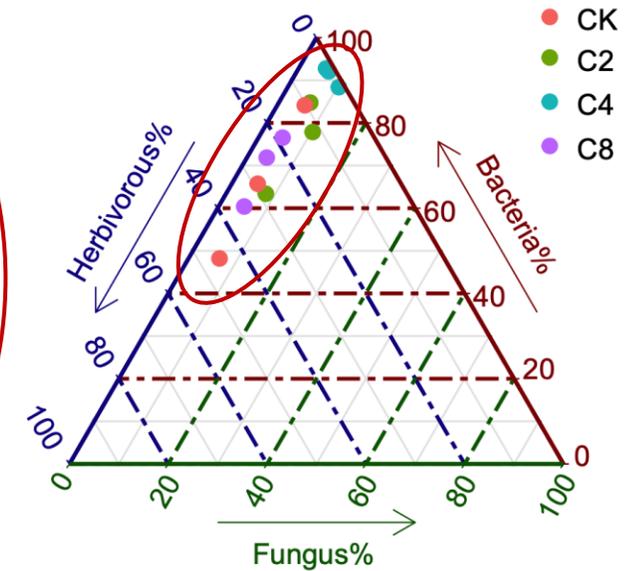
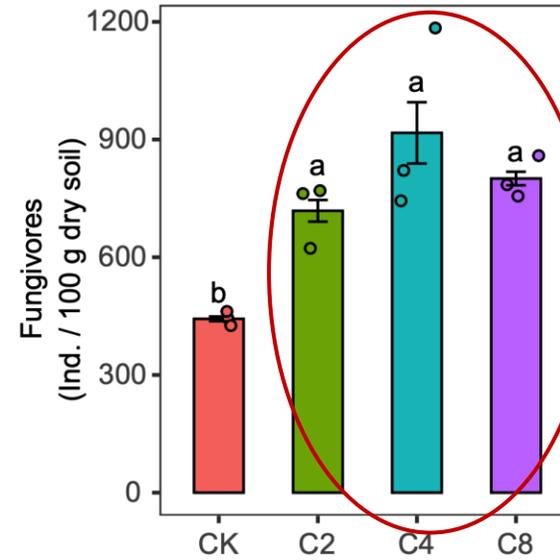
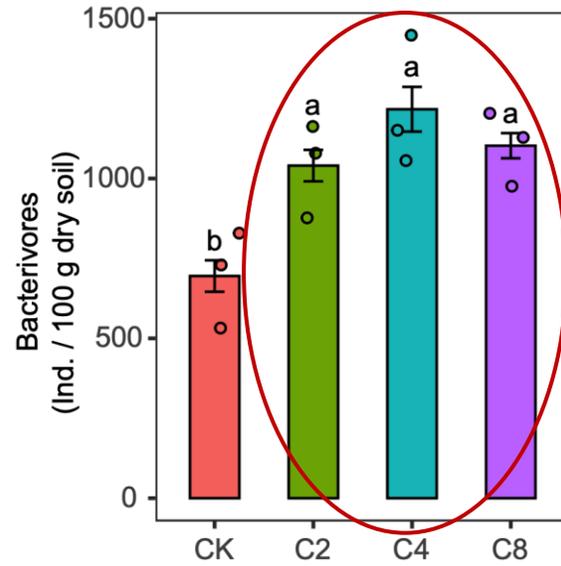


Figure 1. Distribution of trophic groups and life history strategies of soil nematode communities under different cover crop



Case 2: Soil Nematode Ecological Indices

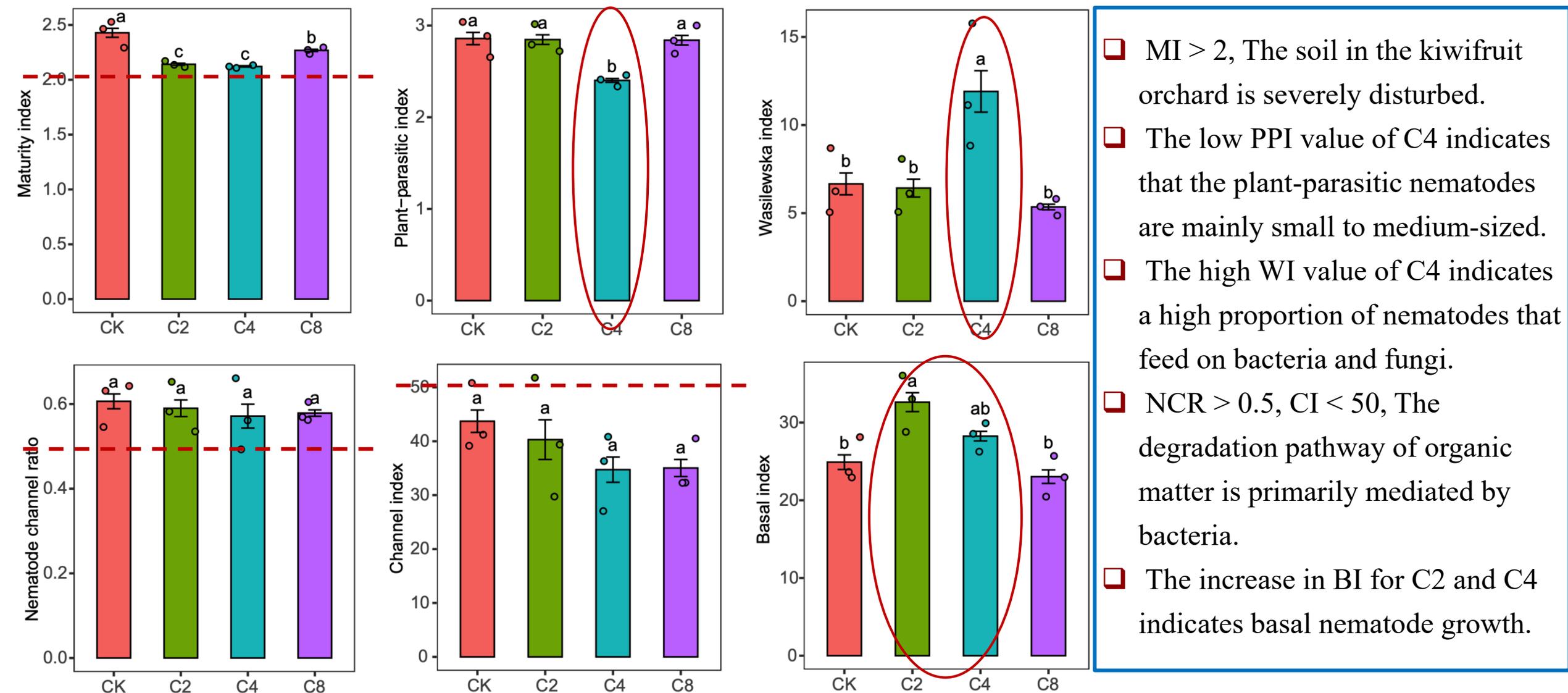
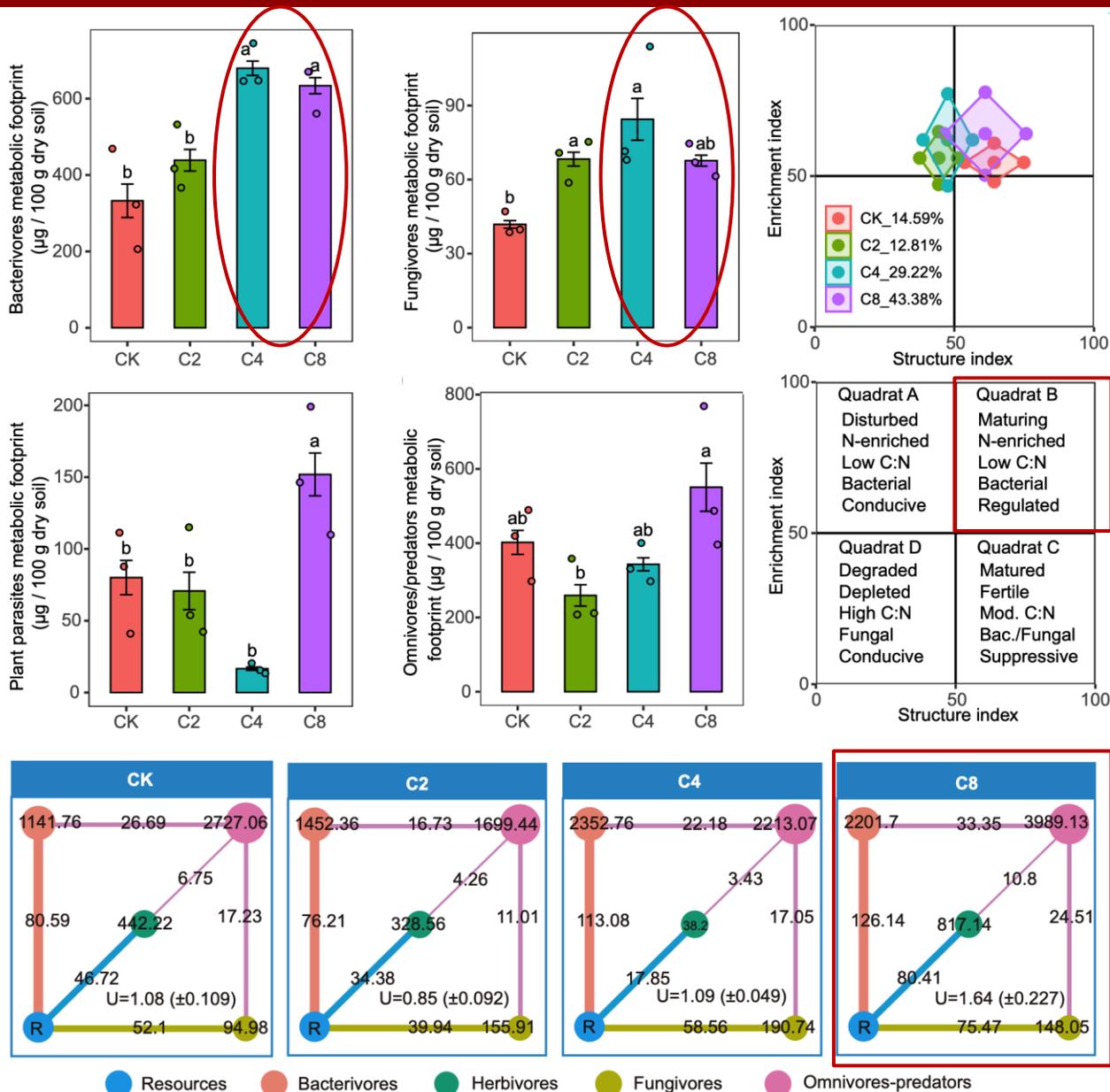


Figure 2. Variations in soil nematode-based indices under different cover crop treatments



Case 3: Metabolic Footprints and Energy Structure

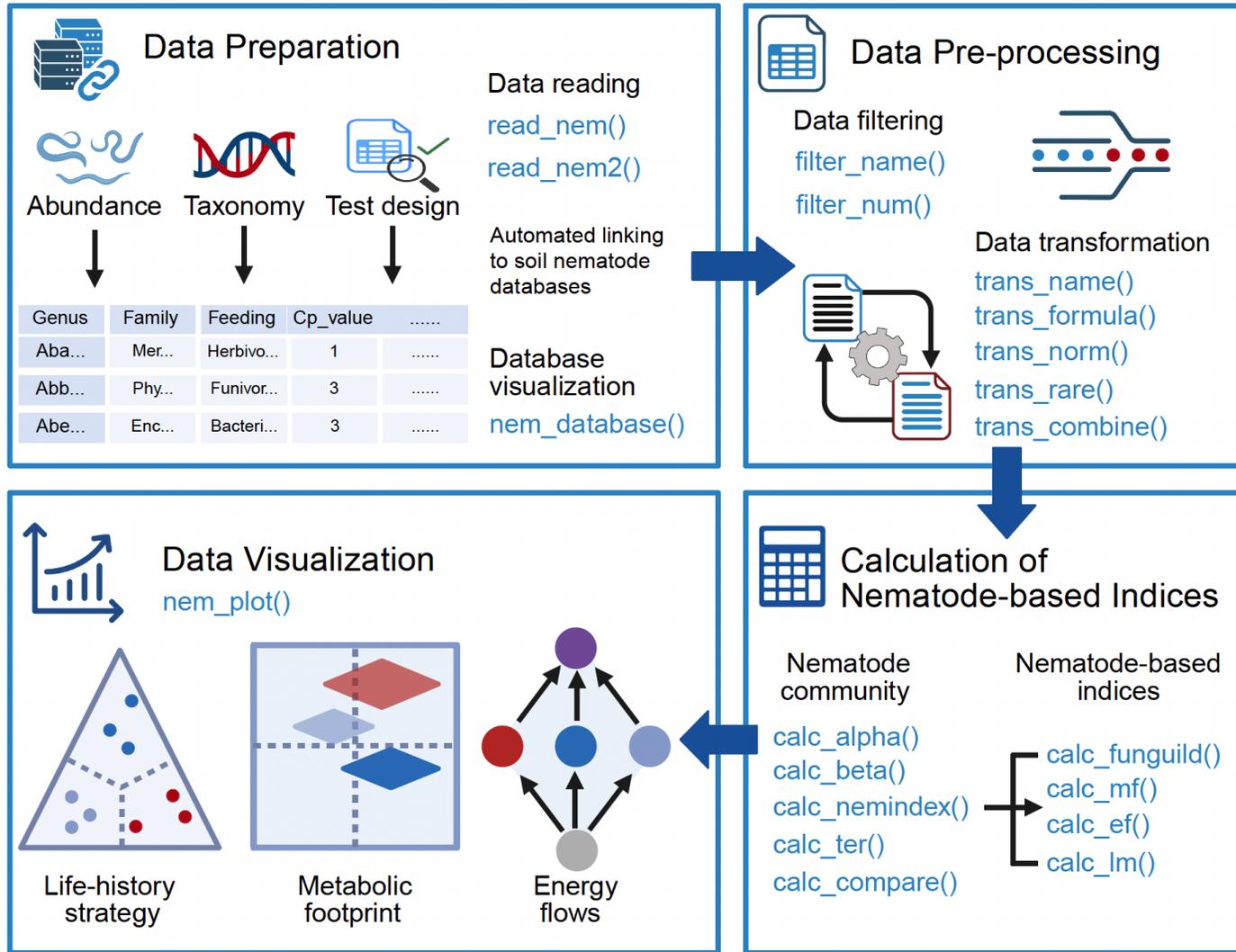


- ❑ C4 and C8 treatments showed significantly higher metabolic footprints for bacterivores and fungivores than CK, indicating enhanced nematode activity and ecological contribution.
- ❑ C8 and CK grouped into faunal analysis Quadrat B, reflecting good nutrient status, low disturbance, and a stable, mature food web.
- ❑ C8 displayed a higher functional metabolic footprint than C2 or C4, suggesting it most effectively promotes predatory omnivore activity.
- ❑ Compared to CK, C8 increased biomass and energy flow across trophic groups, resulting in the most uniform energy structure.

Figure 3. Analysis of metabolic footprint, faunal profiles, and energy structure of soil nematode communities



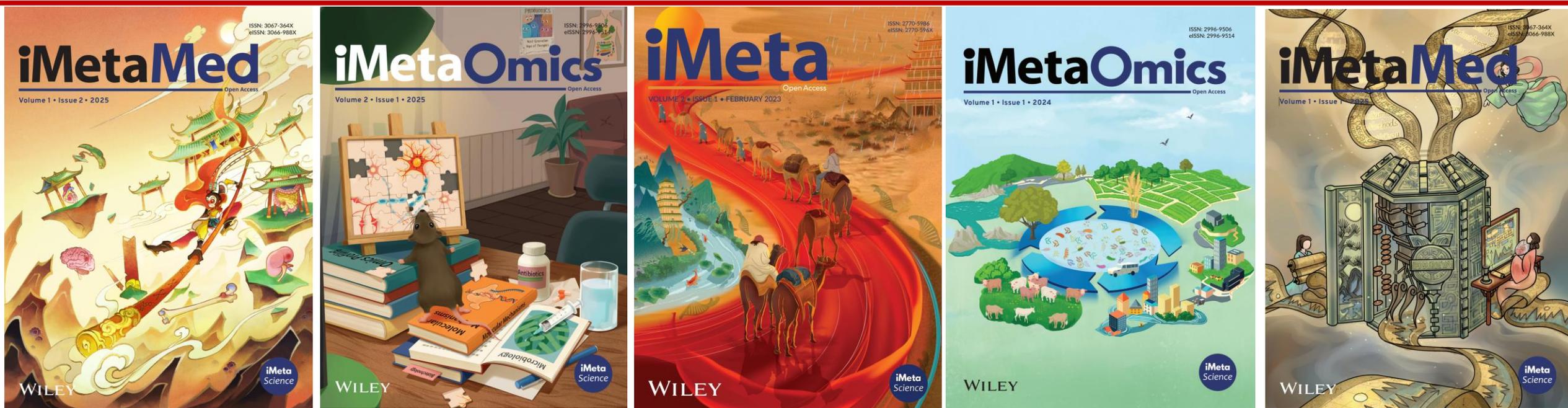
Summary



- *Easynem* is a user-friendly and flexible R package that streamlines the complex analysis of soil nematode communities for evaluating soil health.
- It offers an integrated workflow for data processing, calculation of dozens of ecological indices, and comprehensive visualization of results to generate meaningful ecological insights.

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