Species richness for microbial ecologists

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Goal: Estimate the total number of different taxa



First appearance: Fisher et al. (J Anim Ecol 1943)

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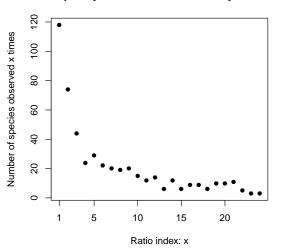
Goal

- Goal: Estimate total (= observed + unobserved) diversity, C
- Sample of n individuals from c distinct classes/species
- Observe f_1 species once singletons f_2 species twice, doubletons f_3 species 3 times... tripletons

 How many species did we observe zero times?

Data structure: Traditional

Frequency abundances for Butterfly dataset

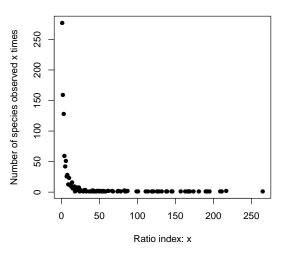


Fisher et al. (J Anim Ecol 1943)

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Data structure: Modern

Frequency abundances for Apples dataset



Walsh et al. (Front Microbiol 2014)

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Richness

Species richness: The total number of different taxa/species/OTUs in the *population*

- Will inherit from the definition your data uses (eg. 97%)
- Key is population

Richness: sample richness

A common, but bad, estimator of the (total, or population) richness is the sample richness

sample richness = number of unique species in your sample

Don't use this! Your sample is not the target of interest!

Richness: Chao1

Another common estimator is the Chao1 "estimator" (or its friends)

$$\begin{aligned} \mathsf{Chao1} &= \mathsf{number} \; \mathsf{of} \; \mathsf{unique} \; \mathsf{species} \; \mathsf{in} \; \mathsf{sample} + \frac{(\# \; \mathsf{singletons})^2}{2 \times \# \; \mathsf{doubletons}} \\ &= c + \frac{f_1^2}{2f_2} \\ \mathsf{s.e.}(\mathsf{Chao1}) &= \sqrt{f_2 \left(\frac{1}{2} \left(\frac{f_1}{f_2}\right)^2 + \left(\frac{f_1}{f_2}\right)^3 + \frac{1}{4} \left(\frac{f_1}{f_2}\right)^4\right)} \end{aligned}$$

We advise against this – too restrictive.

Richness: CatchAll

CatchAll is software for richness estimation

- Fits lots of statistical models and selects best one
- Good for microbial studies in "low-diversity" case
 - ► Homogeneous systems: gut, vaginal...
- Very stable
 - ► Too stable? Underreports errors

Richness: Objective Bayes

The newest distribution of richness software is objective_bayes_*

- Massive improvements in statistical honesty compared to CatchAll
- The same class of models, but better estimation and error quantification
- Just released by Kathryn Barger
 - Low-to-medium-to-medium-high diversity cases?
- Easy to use; available in R

Richness: breakaway

Other good option for richness estimation is breakaway

- Fits different class of statistical models and selects best one
- Good in higher diversity cases
 - ► Non-homogeneous systems: Soil, water...
- A different set of assumptions
 - Pros: More flexible; nests many other cases
 - ► Cons: large standard errors
- Easy to use; available in R

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