

**Anne Chao and Lou Jost. 2012. Coverage-based rarefaction and extrapolation: standardizing samples by completeness rather than size. *Ecology* 93:2533–2547. <http://dx.doi.org/10.1890/11-1952.1>**

APPENDIX B. An unbiased estimator for the expected sample coverage in coverage-based rarefaction.

**Theorem B1:** Under the assumption that sample species frequencies ( $X_1, X_2, \dots, X_S$ ) obey the following multinomial model with cell total  $n$  and cell probabilities ( $p_1, p_2, \dots, p_S$ ):

$$P(X_1 = x_1, \dots, X_S = x_S) = \frac{n!}{x_1! \dots x_S!} p_1^{x_1} p_2^{x_2} \dots p_S^{x_S},$$

the minimum variance unbiased estimator of the expected coverage of a sample with size  $m$ ,

$$E(C_m) = 1 - \sum_{i=1}^S p_i (1 - p_i)^m, \text{ is}$$

$$\hat{C}_m = 1 - \sum_{X_i \geq 1} \frac{X_i}{n} \frac{\binom{n - X_i}{m}}{\binom{n - 1}{m}}, \quad m < n, \quad (\text{B.1})$$

which is also Eq. 4b in the main text. It is readily seen that  $0 \leq \hat{C}_m \leq 1$  for all  $m$ , and

$$1 - \hat{C}_m = \hat{S}_{m+1} - \hat{S}_m,$$

where

$$\hat{S}_m = S_{obs} - \sum_{X_i \geq 1} \frac{\binom{n - X_i}{m}}{\binom{n}{m}}, \quad m < n, \quad (\text{B.2})$$

denotes the species richness estimator in a traditional rarefaction curve for a sample of size  $m < n$ , and  $S_{obs}$  is the number of species observed in the reference sample of size  $n$ .

**Proof:** Under the multinomial assumption, the sample frequency  $X_i$  follows a binomial distribution. Then we have

$$\begin{aligned}
E \left[ \frac{X_i}{n} \frac{\binom{n-X_i}{m}}{\binom{n-1}{m}} \right] &= \sum_{x=1}^n \frac{x}{n} \frac{\binom{n-x}{m}}{\binom{n-1}{m}} \binom{n}{x} p_i^x (1-p_i)^{n-x} \\
&= p_i (1-p_i)^m \sum_{x=1}^n \frac{(n-1-m)!}{(x-1)!(n-x-m)!} p_i^{x-1} (1-p_i)^{n-m-x} \\
&= p_i (1-p_i)^m \sum_{y=0}^{n-1} \binom{n-1-m}{y} p_i^y (1-p_i)^{n-1-m-y} \\
&= p_i (1-p_i)^m.
\end{aligned}$$

From the Rao-Blackwell and Lehmann-Scheffé Theorems (e.g., Casella and Berger 2002, p. 347 and p. 369), the estimator in Eq. B.1 is the unique minimum variance unbiased estimator of the expected sample coverage. Also,  $1 - \hat{C}_m = \hat{S}_{m+1} - \hat{S}_m$  follows directly from the following equation:

$$\frac{\binom{n-X_i}{m}}{\binom{n-1}{m}} - \frac{\binom{n-X_i}{m+1}}{\binom{n-1}{m+1}} = \frac{X_i}{n} \frac{\binom{n-X_i}{m}}{\binom{n-1}{m}}.$$

**Theorem B2:** We have the following monotonic properties:

- (1) The four functions  $E(S_m)$ ,  $E(C_m)$ ,  $\hat{S}_m$  and  $\hat{C}_m$  are all non-decreasing function of sample size  $m$ ;
- (2)  $E(S_m)$  is a non-decreasing function of  $E(C_m)$ ;
- (3)  $\hat{S}_m$  is a non-decreasing function of  $\hat{C}_m$ .

Proof: For any  $X_i \geq 1$ , direct computation leads to

$$\frac{\binom{n-X_i}{m+1}}{\binom{n-1}{m+1}} \leq \frac{\binom{n-X_i}{m}}{\binom{n-1}{m}}.$$

This implies  $\hat{C}_{m+1} \geq \hat{C}_m$ . Since  $\hat{S}_{m+1} - \hat{S}_m = 1 - \hat{C}_m \geq 0$ , we obtain  $\hat{S}_{m+1} \geq \hat{S}_m$ . The other monotonic properties are readily seen.

#### LITERATURE CITED

Casella G., and R. L. Berger. 2002. Statistical Inference, Second Edition. Duxbury, Pacific Grove.