¹ Appendix A.

² Analytic estimators for rarefaction (interpolation) of Hill numbers (${}^{q}\hat{H}$) of order q = 0 (first row

³ of equations), q = 1 (second row), and q = 2 (third row), given a reference sample (Chao et al.

- ⁴ 2014). The last row gives equations for sample completeness (coverage, \hat{C}) as a function of
- sample size. In abundance-based rarefaction: n is the reference sample; m is defined as the
- ⁶ subsample size for rarefying community; S_{obs} is the observed species richness in n; X_i is a vector
- ⁷ of length S_{obs} , the elements of which are the observed abundances; k is the vector of sampling
- ⁸ units, from 1 to m; and $\hat{f}_k(m)$ is an unbiased estimator of the number of species detected in

• exactly *m* sampling units, where
$$\hat{f}_k(m) = \sum_{X_i \ge k} \frac{\begin{pmatrix} X_i \\ k \end{pmatrix} \begin{pmatrix} n - X_i \\ m - k \end{pmatrix}}{\begin{pmatrix} n \\ m \end{pmatrix}}$$
, provided that $m < n$, and $k \ge 1$.

In sample-based rarefaction: *T* is the reference sample; *t* is defined as the subsample size for rarefying community; S_{obs} is the observed species richness in *T*; Y_i is the matrix of incidence data, with i = 1 to S_{obs} rows and j = 1 to *T* columns; *k* is the vector of sampling units, from 1 to *t*; \hat{U}_t is an unbiased estimator of the expected total number of incidences (*U*) in *t* sampling units,

¹⁴ $\hat{U}_t = t \times U/T$; and $\hat{Q}_k(m)$ is an unbiased estimator of the number of species detected in exactly *k*

sampling units, where
$$\hat{Q}_k(t) = \sum_{Y_i \ge k} \frac{\begin{pmatrix} Y_i \\ k \end{pmatrix} \begin{pmatrix} T - Y_i \\ t - k \end{pmatrix}}{\begin{pmatrix} T \\ t \end{pmatrix}}$$
, provided that $t < T$, and $t \ge 1$. For the

¹⁶ reference sample, the observed Hill number of order q is ${}^{q}\hat{D}(n) = {}^{q}D_{obs} = \left[\sum_{Xi \ge 1} (X_{i}/n)^{q}\right]^{1/(1-q)}$ in ¹⁷ abundance-based rarefaction, and ${}^{q}\hat{\triangle}(T) = {}^{q}\triangle_{obs} = \left[\sum_{k=1}^{t} (k/U)^{q} \times Q_{k}\right]^{1/(1-q)}$ in incidence-based

rarefaction. The coverage of the reference sample is estimated by $\hat{C}(n) = 1 - \frac{f_1}{n} \left[\frac{(n-1) \times f_1}{(n-1) \times f_1 + 2 \times f_2} \right]$ in abundance-based rarefaction, ds and $\hat{C}(T) = 1 - \frac{Q_1}{T} \left[\frac{(T-1) \times Q_1}{(T-1) \times Q_1 + 2 \times Q_2} \right]$ in incidence-based rarefaction.

| 21 | | | |
|----|----------|--|---|
| | | Abundance-based $(m < n)$ | Incidence-based $(t < T)$ |
| | q = 0 | ${}^{0}\hat{H}(m) = \sum_{k=1}^{m} \hat{f}_{k}(m) = S_{obs} - \sum_{Xi \ge 1} \frac{\begin{pmatrix} n - X_{i} \\ m \end{pmatrix}}{\begin{pmatrix} n \\ m \end{pmatrix}}$ | ${}^{0}\hat{H}(t) = S_{obs} - \sum_{Yi \ge 1} \frac{\begin{pmatrix} T - Y_i \\ t \end{pmatrix}}{\begin{pmatrix} T \\ t \end{pmatrix}}$ |
| 22 | q = 1 | ${}^{1}\hat{H}(m) = \exp\left(\sum_{k=1}^{m} \left(-\frac{k}{m} \times \log\frac{k}{m}\right) \times \hat{f}_{k}(m)\right)$ | ${}^{1}\hat{H}(t) = exp\left(\sum_{k=1}^{t} \left(-\frac{k}{\hat{U}_{t}} \times \log \frac{k}{\hat{U}_{t}}\right) \times \hat{Q}_{k}(t)\right)$ |
| | q = 2 | ${}^{2}\hat{H}(m) = rac{1}{\sum\limits_{k=1}^{m} \left(rac{k}{m} ight)^{2} 	imes \hat{f}_{k}(m)}$ | ${}^{2}\hat{H}(t)=rac{1}{\sum\limits_{k=1}^{m}\left(rac{k}{\hat{U}_{t}} ight)^{2}	imes\hat{\mathcal{Q}_{k}}(t)}$ |
| | Coverage | $\hat{C}(m) = 1 - \sum_{X_i \ge 1}^{m} \frac{X_i \begin{pmatrix} n - X_i \\ m \end{pmatrix}}{n \begin{pmatrix} n - 1 \\ m \end{pmatrix}}$ | $\hat{C}(t) = 1 - \sum_{Y_i \ge 1}^{m} \frac{Y_i \begin{pmatrix} T - Y_i \\ t \end{pmatrix}}{n \begin{pmatrix} T - 1 \\ t \end{pmatrix}}$ |