The introduction of species abundance distribution

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The following six Species abundance distribution (SAD) models were considered:

broken-stick, niche-preemption, log-normal, Zipf, Zipf-Mandelbrot, and neutral-theory models

(Table 1). Further details and comments of other SAD models are described by McGill et al. (2007)

and Wilson (1991).

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Model	Equation		Reference	
Broken-stick	$\hat{a}_r = \frac{N}{S} \sum_{k=r}^{S} \frac{1}{k}$	(1)	MacArthur (1957)	
Niche-preemption	$\hat{a}_r = N\alpha(1-a)^{r-1}$	(2)	Motomura (1932)	
Log-normal	$\hat{a}_r = \exp[\log(u) + \log(\sigma)\Phi]$	(3)	Preston (1948)	
Zipf	$\hat{a}_r = N \hat{p}_1 r^{\gamma}$	(4)	Frontier (1987)	
Zipf-Mandelbrot	$\hat{a}_r = Nc(r+\beta)^{\gamma}$	(5)		
Neutral-theory	$\phi_n = \theta \frac{J!}{n! (J-n)!} \frac{\Gamma(\gamma)}{\Gamma(J+\gamma)} \int_0^{\gamma} \frac{\Gamma(n+y)}{\Gamma(1+y)} \frac{\Gamma(J-n+\gamma-y)}{\Gamma(\gamma-y)} \exp(-y\theta/\gamma) dy$	(6)	Hubbell (2001)	

Note: \hat{a}_r is the expected abundance of species at rank r, *S* is the number of species, *N* is the number of individuals, Φ is a standard normal function, \hat{p}_1 is the estimated proportion of the most abundant species, and $\alpha, \sigma, \gamma, \beta$ and *c* are the estimated parameters in each model. In neutral-theory model, where $\Gamma(z) = \int_0^\infty t^{z-1} e^{-t} dt$ which is equal to (z-1)!, for integer z and $\gamma = \frac{m(J-1)}{1-m}$, θ is fundamental diversity number, *m* is migration rate.

Broken-stick model: This model was first proposed by MacArthur (1957). Its analogy of placing *s*-points randomly on a line of unit length and simultaneously breaking it at those points into *s* lengths can be rephrased as a group of *s* series. The lengths of the segments represent the "niche sizes" of the species. According to the model, the expected size of the *r*th species, and \hat{a}_r , the expected abundance of species of species at rank *r*, are shown in equation (1) in Table 1. The mathematical proof of this model can be found in Pielou (1975).

Niche-preemption model This model was proposed by (Motomura, 1932) and assumes that the percentage of the total niche occupied by the first species is α , the second species occupied a percentage α of the reminder, $\alpha(1 - \alpha)$, and so on. The expected abundance for the *r*th species is equation (2) in Table 1.

Log-normal model A log-normal distribution is defined as a distribution whose variate conforms to the normal laws of probability. For SADs, the log-normal distribution characterizes a sample with relatively low abundance or very rare species (Matthews and Whittaker, 2014). Preston (1948) introduced the log-normal SAD by demonstrating a good fit to a large number of data sets covering a number of different communities. See equation (3) in Table 1.

Zipf and Zipf-Mandelbrot model The Mandelbrot model was originally developed for information systems, assessing the cost of information (Frontier, 1985). In plant communities, the presence of a species can be seen as dependent on previous physical conditions and previous species presences – these are the costs. Pioneer species have a low cost, requiring few prior conditions. Late successional species have a high cost, viz. the energy, time, and organization of the ecosystem required before they can invade. On this basis they will be rare (Frontier, 1987). These differences between species give a Zipf or Zipf-Mandelbrot distribution, equations (4) and (5) in Table 1, respectively. The assumption is that a species is very likely to invade once its necessary conditions are met (Wilson, 1991).

Neutral-theory model Hubbell (2001) noted that the relative abundance of species within – and the species diversity of – a community can be explained through neutral drift of individual species abundances. The model contends that the number of individuals in a metacommunity is constant, that is, all available resources in the community are saturated. This is the zero-sum assumption: if an individual dies and a portion of the resource becomes available, it will be immediately taken up by a new individual, and the community size remains constant. See equation

(6) in Table 1.

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