

Testing the Spatial Predictions of the Maximum Entropy Theory of Ecology

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Introduction

Many macroecological patterns may simply reflect emergent statistical phenomena rather than specific ecological processes.

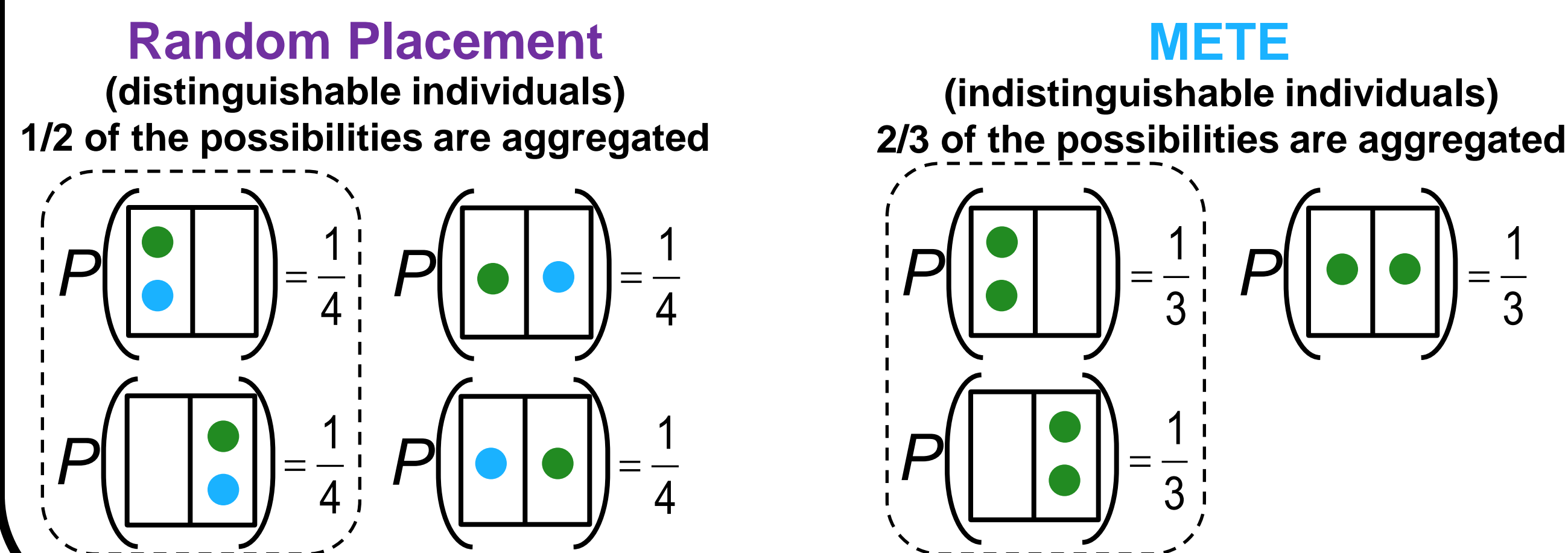
The Maximum Entropy Theory of Ecology (METE) takes advantage of this fact to unify community patterns through four state variables: 1) number of species, 2) number of individuals, 3) energy, 4) area.

METE ignores specific ecological processes operating below the level of the state variables. METE asserts that ecological patterns should be characterized by the most likely state of the system that maintains empirically observed constraints on the state variables.

We develop and test the spatially explicit predictions of METE using the species-area and distance-decay relationships across a diverse set of plant communities.

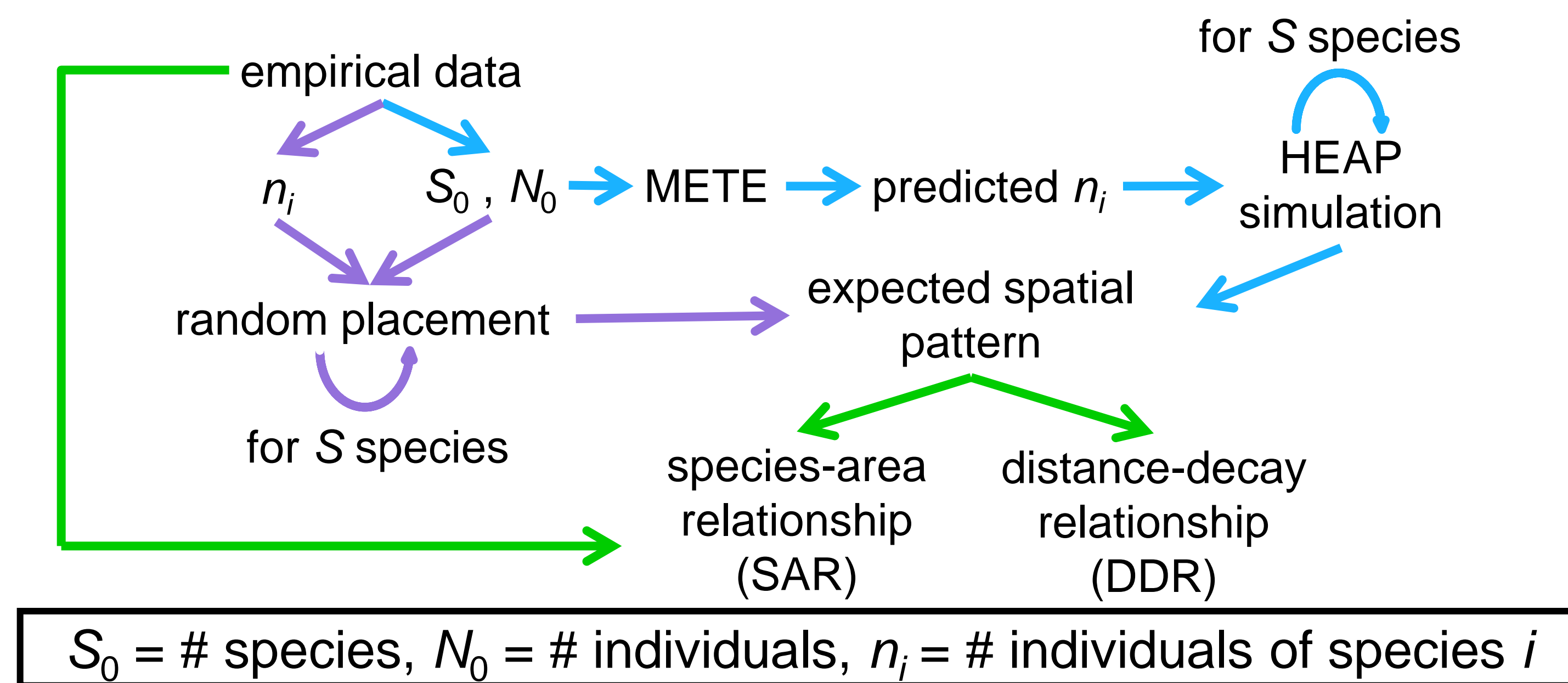
Two Models of Spatial Pattern

- Random placement assumes that individuals are **distinguishable**
- METE assumes that individuals are **indistinguishable**
- Hypothesis of Equal Allocation Probability (HEAP)**
 - all unique spatial arrangements of individuals are equally likely
- Under HEAP, if two individuals are placed in either the left or right half of a quadrat, there is a higher probability that individuals will be aggregated if they are indistinguishable



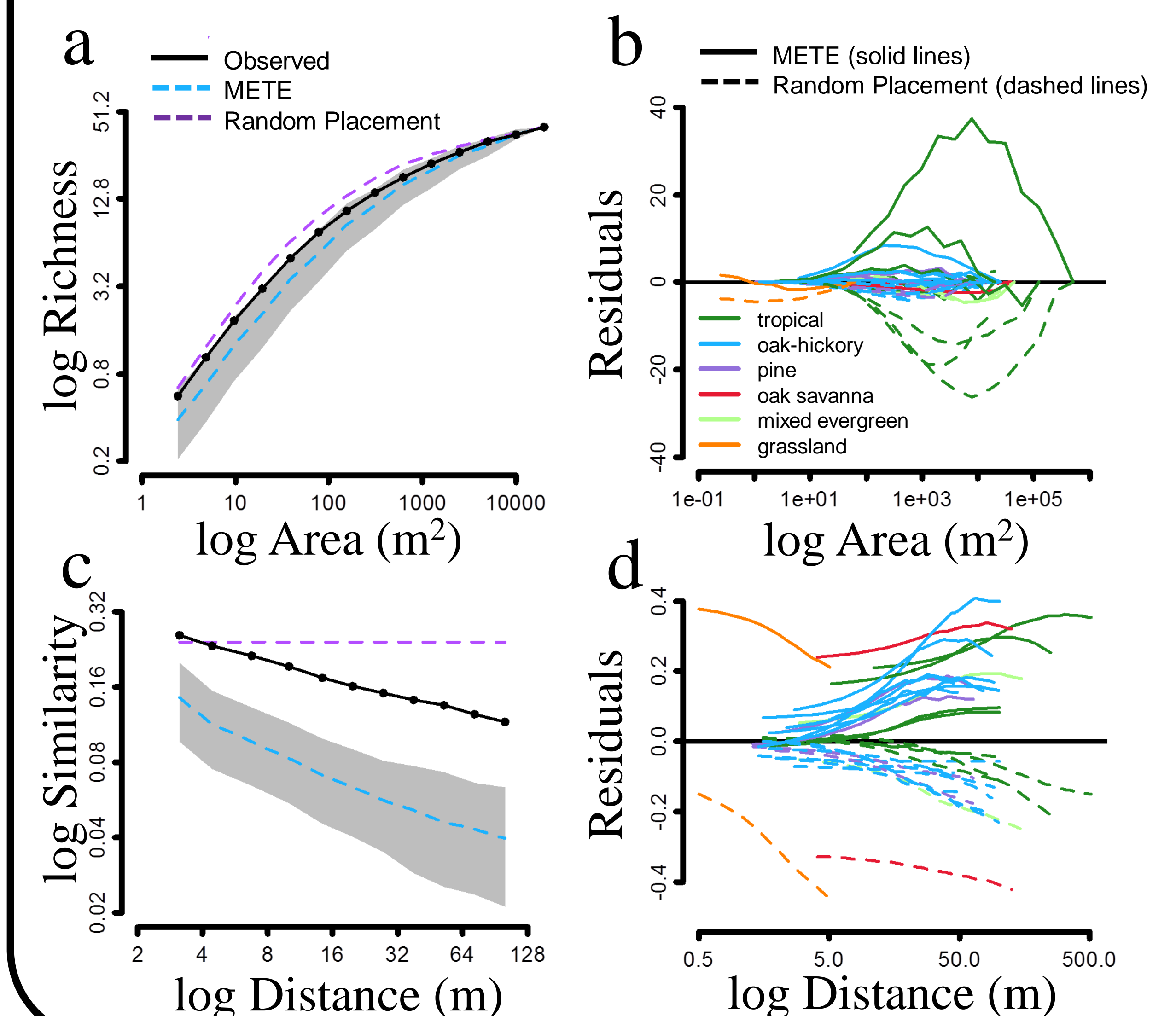
Objectives

- Develop METE's spatially explicit predictions
- Examine the degree of fit to empirical data
- Compare METE's performance to the hypothesis of random placement



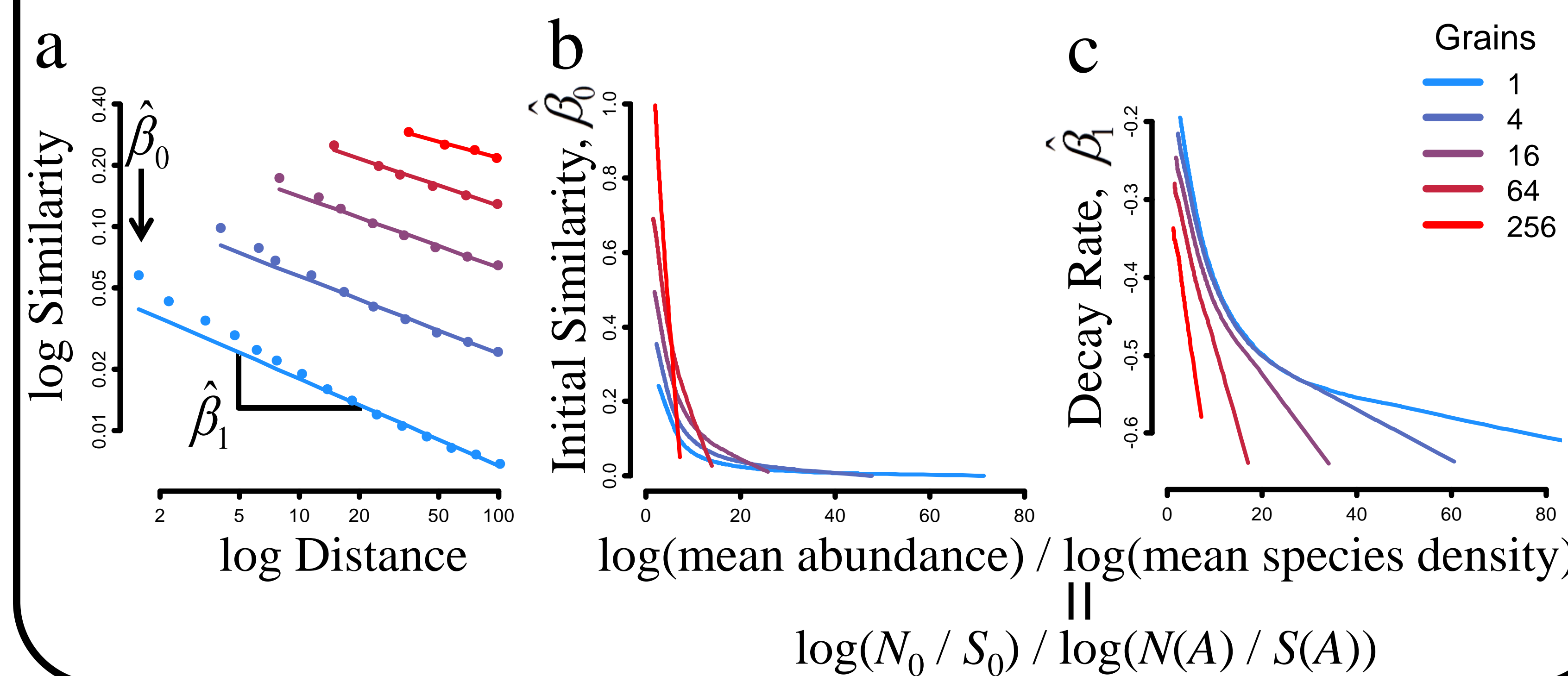
Empirical Results

Panels (a, c) display typical observed patterns of the SAR and DDR respectively, (b, d) display the residuals of the SAR and DDR across all datasets with respect to the METE and random placement models. **Overall the random placement model outperformed METE for both the SAR and DDR.**



METE Distance Decay

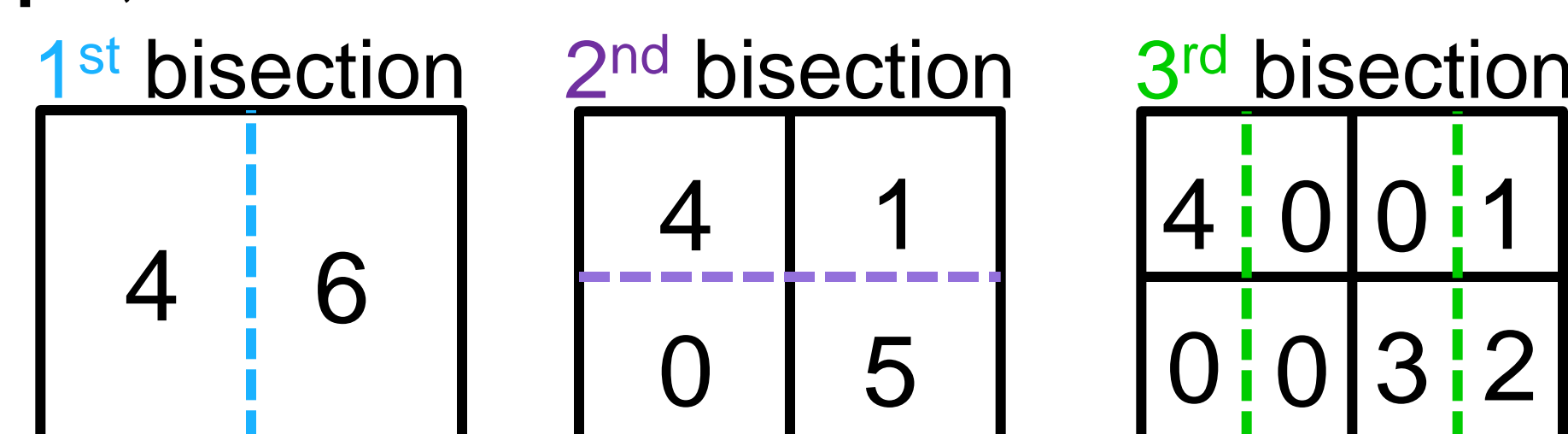
The METE DDR of Sorensen's Similarity index, are well described by a power model (a). The intercept $\hat{\beta}_0$ and slope $\hat{\beta}_1$ of the model are controlled by a ratio of N and S (b, c).



Spatial Simulation Approach

We developed the spatially explicit predictions for METE by iteratively bisecting an area and applying HEAP until the spatial grain of interest was achieved.

For example, with 10 individuals and 3 bisections



At each bisection the individuals are randomly placed on either side of the bisection line, this can quickly generate spatially aggregated patterns of individuals.

Empirical Datasets

Habitat type	# sites	Extent (ha)	S	N
tropical forest	4	2.00 – 50	124 – 301	4326 – 205096
oak-hickory forest	7	0.5 – 2.56	17 – 48	669 – 8887
old field pine forest	2	0.85 – 1	36 – 41	2139 – 2584
oak savanna	1	4	7	7625
mixed evergreen forest	1	4.5	31	5885
serpentine grassland	1	0.0064	24	37182
Summary	16	0.0064 – 50	7 – 301	669 – 205096

Conclusions

We have uncovered that METE predicts a log-log linear pattern of distance decay. The characteristics of the METE DDR are controlled by a ratio of the state variables. METE appeared to capture the functional form of the SAR and DDR, but it fit the data quite poorly. Although overall random placement performed better than METE, it appears that the empirical data may be best described by a middle ground between these two models. Future models should consider a model of intra-specific aggregation that is not random but is less aggregated than METE. The state variable approach, upon which these models are built, provides a simple but powerful framework for unifying community patterns.

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