

# A Phylogenetic Perspective on the Individual Species-Area Relationship in Temperate and Tropical Tree Community

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Species coexistence

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niche-based theory (e.g. Tilman 1982)

neutral theory (e.g. Hubbell 2001; Chave 2004)
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Spatial distribution

leave a spatial signature (Hubbell et al. 2001)

Species-area relationships (SARs)



# How individual species structure diversity in tropical forests

Thorsten Wiegand\*\*, C. V. Savitri Gunatilleke\*, I. A. U. Nimal Gunatilleke\*, and Andreas Huth\*

**Species-area relationships (SARs)** 





# How individual species structure diversity in tropical forests

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**Species-area relationships (SARs)** 

**■ Individual species-area relationships (ISARs)** 

Role of species

accumulator repeller neutral species

Species area relationships (SARs)
Individual species area relationships (ISARs)

**Species area relationships (SARs)** 

Individual species area relationships (ISARs)



Individual phylogentic area relationships (IPARs)

- Do ISARs change predictably across latitude?
- Is the phylogenetic diversity in the neighborhood of species accumulators and repellers higher or lower than that expected given the observed species richness?
- What is the distribution of accumulator, repeller and neutral species on the phylogenetic tree?

Study sites

FDP	Country	Forest Type	Latitude	Area (ha)	Species Richness
Ailaoshan	China	Subtropical Moist Forest	24.32 N	6	76
BCI	Panama	Tropical Moist Forest	9.154 N	50	299
Edoro – 1	Congo	Tropical Rain Forest	1.437 N	10	315
Edoro – 2	Congo	Tropical Rain Forest	1.437 N	10	326
Korup	Cameroon	Tropical Rain Forest	5.074 N	50	494
Lenda – 1	Congo	Tropical Rain Forest	1.437 N	10	349
Lenda – 2	Congo	Tropical Rain Forest	1.437 N	10	300
Wabikon Lake	U.S.A.	Temperate Deciduous Forest	45.551 N	25	36
Xishuangbanan	China	Tropical Rain Forest	21.612 N	20	468

- Study sites
- Species selection

Totally 2,728 species, 1,020,699 individuals

- all species
- species that abundant ≥ 70 and DBH ≥ 1 cm

- Study sites
- Phylogenetic tree reconstruction
  - Phylomatic (Webb and Donoghue 2005)
  - Phylocom (Webb et al. 2008)

- Study sites
- Phylogenetic tree reconstruction
- Individual species-area relationships
  - ISAR (Wiegand et al. 2007)
  - heterogeneous null model



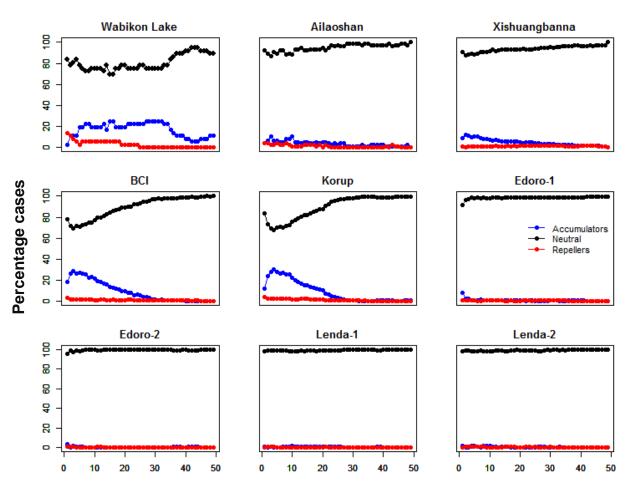
- Study sites
- Phylogenetic tree reconstruction
- Individual species-area relationships
- Individual phylogenetic-area relationships
  - phylogenetic diversity metric
  - tip shuffle null model

- Study sites
- Phylogenetic tree reconstruction
- Individual species-area relationships
- Individual phylogenetic-area relationships
- Phylogenetic distribution of accumulator, repeller and neutral species
  - NRI & NTI

#### **Results and Discussion**

- Individual species-area relationships
- Individual phylogenetic-area relationships
- Phylogenetic distribution of accumulator, repeller and neutral species

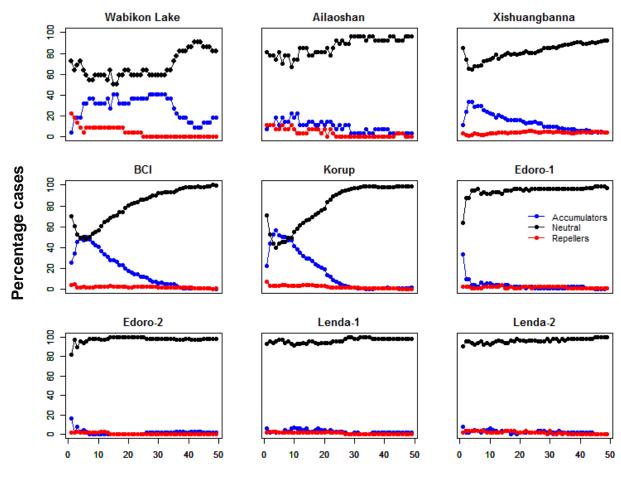
Individualspecies-arearelationshipsall species



Distance r from target tree [m]

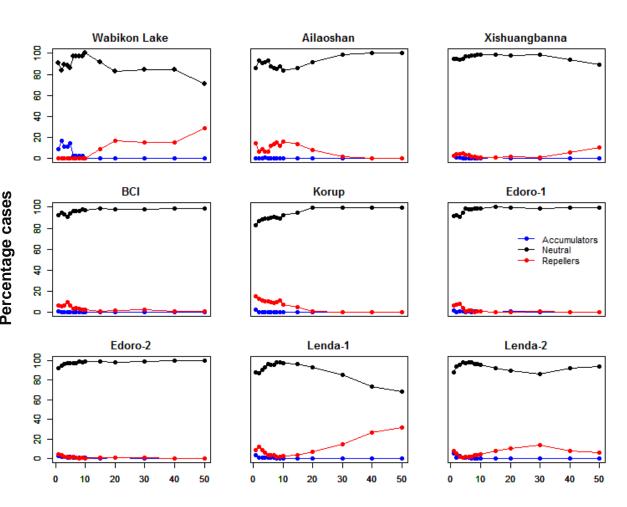
Individualspecies-arearelationships

species that abundant ≥ 70 and DBH ≥ 1 cm



Distance r from target tree [m]

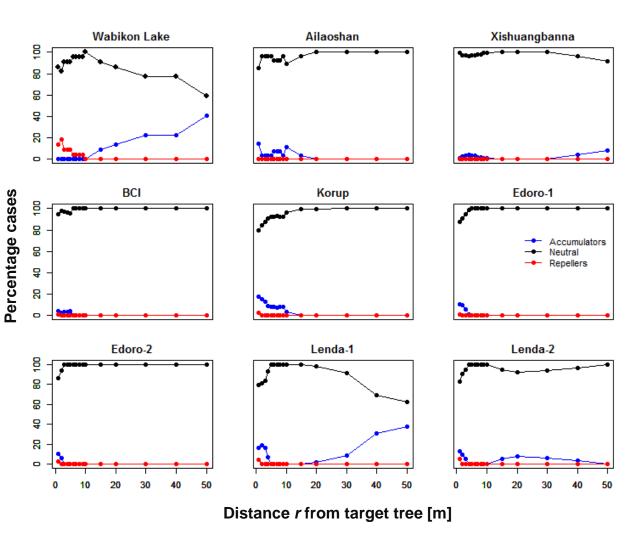
Individual
 phylogentic-area
 relationships
 all species



Distance r from target tree [m]

Individualphylogentic-arearelationships

species that
abundant ≥ 70
and DBH ≥ 1 cm



#### **Conclusions**

- Non-neutral processes (competition and facilitation) may leave a detectable signature at small-scales spatial patterns of species diversity but result in stochastic patterns at larger-scales.
- Importance of past evolutionary history in dictating the ecological interactions we presently observe.

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