

Differences in the spatial structure of the primary and secondary tropical rain forests

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Spatial analyses and TRFs

We expect that during stand aging

- patterns start from clumped (aggregated) and change to random or regular
- decreasing autocorrelation of plant sizes

Focus on the spatial patterns in TRFs

- the first analyses by Greig-Smith in Trinidad in early 50s
- strong aggregation observed (Hubbell 1979; Condit et al. 2000) explained due to environmental heterogeneity (Bagchi et al. 2011)
- after separating environmental heterogeneity interactions of individuals should remain

Diversity of TRFs and Questions

- the most of species is diversity neutral (Wiegand et al. 2007)
- segregation and aggregation decrease with species richness (Wiegand et al. 2012)

Our questions

Using homogeneous TRF plots

- what spatial differences and which species interactions we can observe in different successional stages of TRF?
- what are spatial diversity patterns of successional stages of TRF?

Study sites

- tropical rain forest (TRF) in Papua New Guinea
 - two homogeneous 1 ha plots
 - 100-200 m above sea level
 - annual rainfall of 3500 mm
 - annual mean monthly temperature around 26.5 C
- **primary TRF** "late succession"
minimally >60 years no human disturbance
- **secondary TRF** "early succession"
abandoned garden for a decade
- 2456 trees with DBH ≥ 5 cm :
position, species identity, DBH, height

Border of primary TRF



Myristicaceae, Horsfieldia basifissa, Teijsmanniodendron bogoriense 5/26

Border of secondary TRF



Euphorbiaceae, Macaranga tanarius, Ficus variegata

Study sites - field work *F. nodosa*



Study sites - common species

Horsfieldia basifissa (Primary), *Ficus pungens* (Secondary)



Marked points pattern analyses

Individuals pattern

- **pair** correlation function

Marks pattern

- **mark** correlation function
- **mark variogram** (mark autocorrelation)
- **spatial diversity** function ("K-function + SAR")
 - species number and Simpson diversity along spatial scale

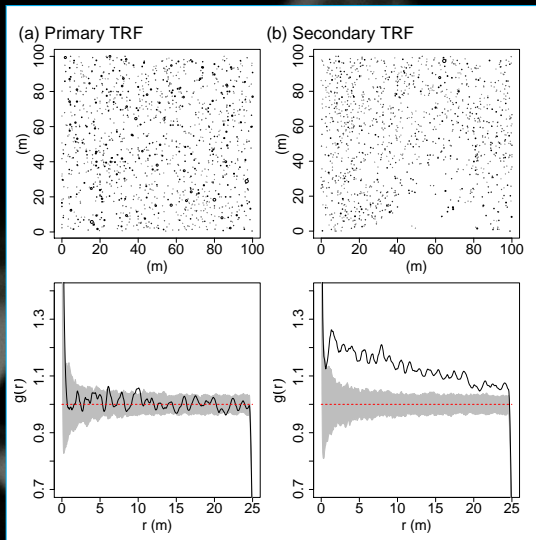
Null models

- **no-marks**: random and inhomogeneous positions
- **marks**: random relabelling and species shifting

Comparison in numbers

Characteristics	Primary TRF	Secondary TRF
# individuals	1255	1201
Basal area (m ² /ha)	29.9	13.6
# species	198	88
# species with 1 individual	64	36
# species with ≥ 20 individuals	12	12
- # these individuals	562	974
# shared species	45	45
- # these individuals	579	821

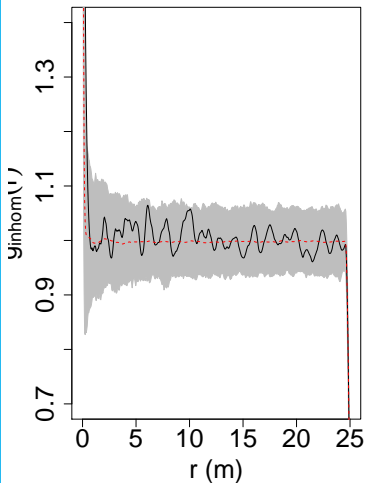
Plots and their pair corr. functions



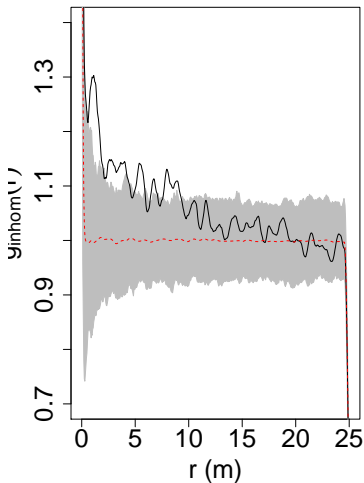
gray area = null model, solid line = observed pattern

Inhomogeneous pair corr. functions

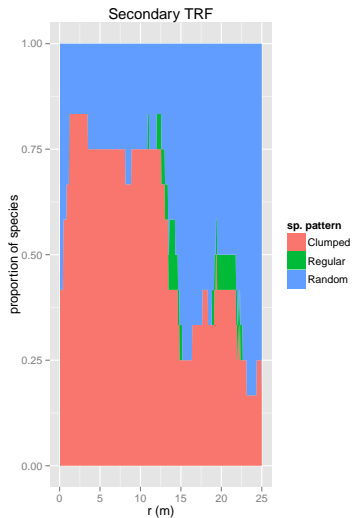
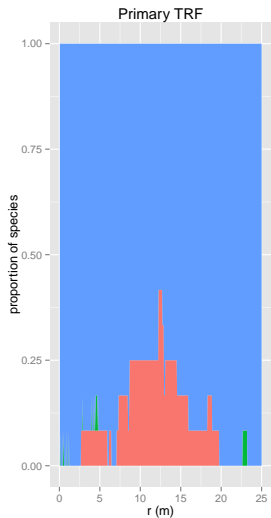
(a) Primary TRF



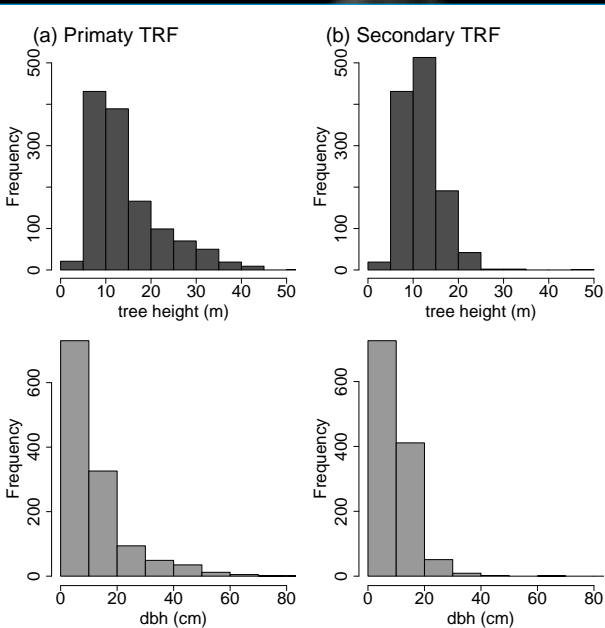
(b) Secondary TRF



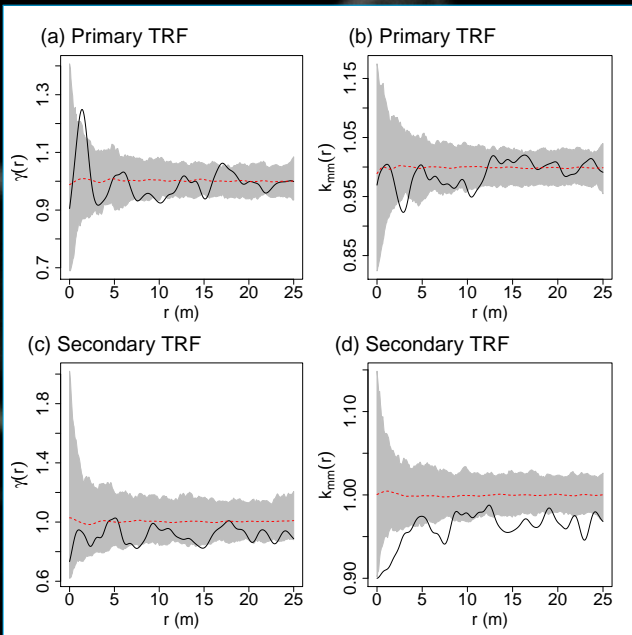
Proportions of patterns of 12 most common species



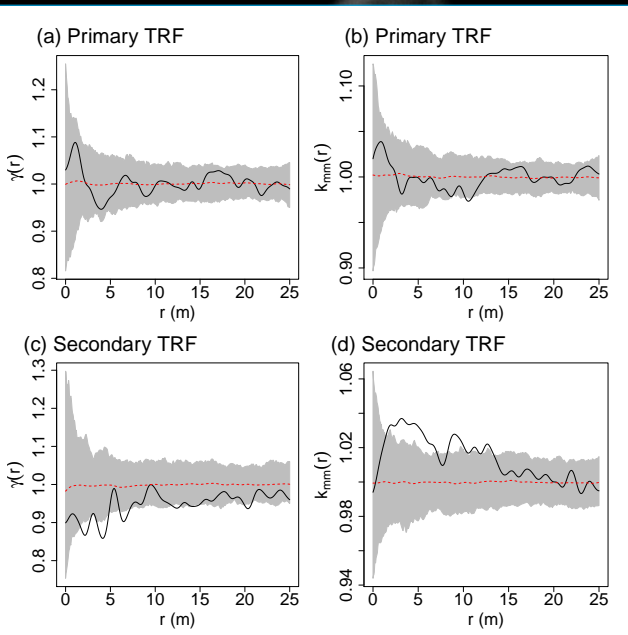
Height and DBH histograms



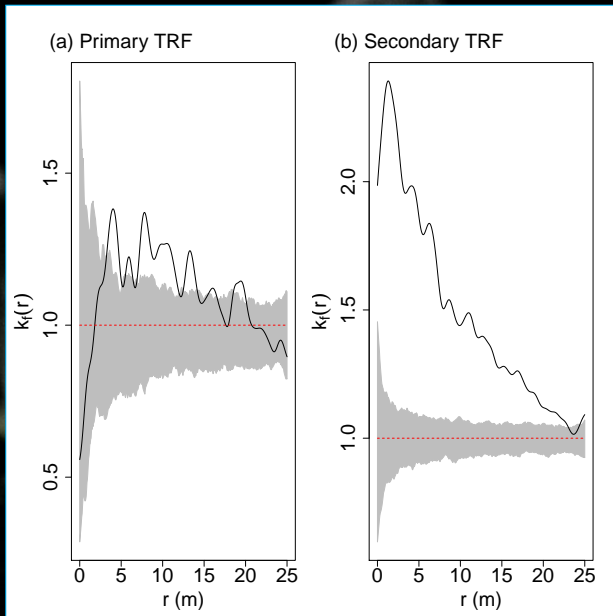
$\gamma(r)$ - vario., $\kappa(r)$ - corr. marks DBH



$\gamma(r)$ - vario., $\kappa(r)$ - corr. marks height

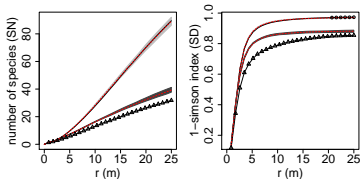


$\kappa(r)$ mark corr. of species identity

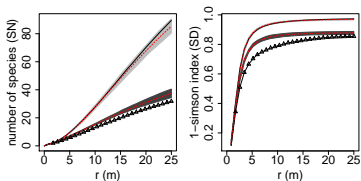


Species number and diversity

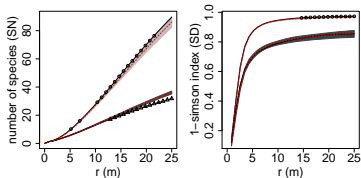
random labelling



inhom. positions



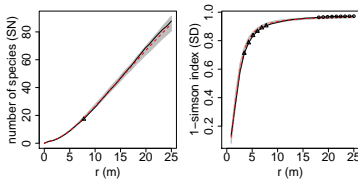
shifting of species



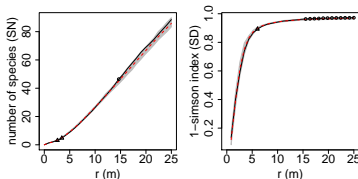
light gray = primary TRF, dark gray = secondary TRF

Spatial div. - Primary TRF shifting

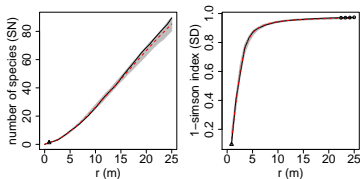
Horsfieldia basifissa
(122 inds.)



Teijsmanniodendron bogoriense (66 inds.)

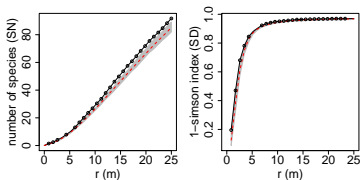


Gymnacranthera paniculata (61 inds.)

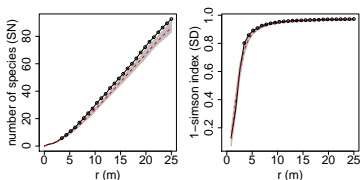


Spatial div. - Primary TRF shifting

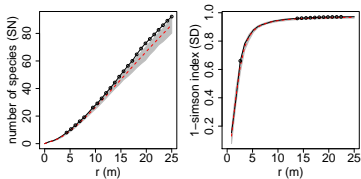
Pometia pinnata (53
inds.)



*Pimelodendron am-
boinicum* (50 inds.)

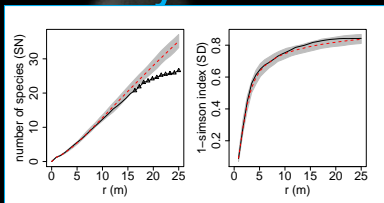


*Mastixiodendron
pachyclados* (50
inds.)

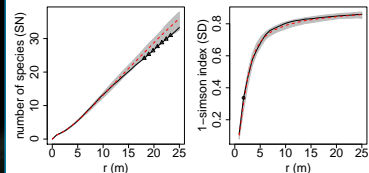


Spatial div. - Secondary TRF shifting

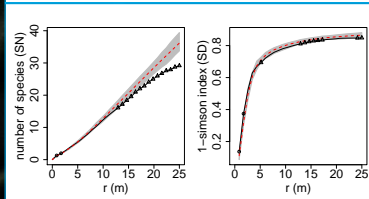
Macaranga tanarius
(275 inds.)



Ficus variegata (220
inds.)

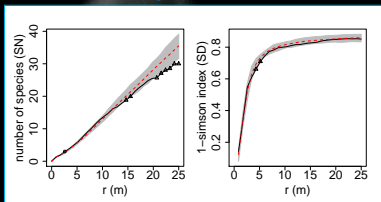


*Trichospermum
pleiostigma* (164
inds.)

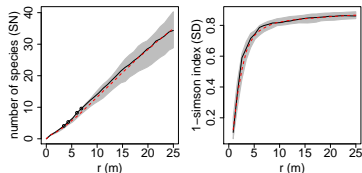


Spatial div. - Secondary TRF shifting

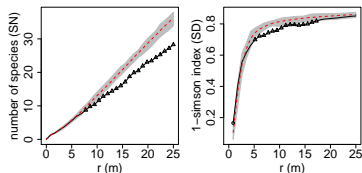
Macaranga aleuritoides (81 inds.)



Vitex cofassus (47 inds.)



Trema orientalis (34 inds.)



Spatial patterns, DBH and height

–conclusions

- in the homogeneous primary TRF, there was mostly random pattern of all and within the most common species individuals
- early succession stage, the secondary TRF, was clumped (all and the most common species individuals), even if inhomogeneity was filtered out
- distribution of DBH and height
 - random in the primary TRF
 - clumps of thin and clumps of high individuals in the secondary TRF

Spatial diversity – conclusions

- the results out of confidence envelopes mostly at the larger scales
- in the secondary TRF, clumps of conspecific neighbours followed by low spatial diversity mostly due to dominant species functioning as "diversity repellers"
- in the primary TRF, less negative or more positive inter-specific interactions than intra-specific interactions, some of dominant species were "diversity accumulators"

Future work and Questions

Spatial investigation of

- plant traits
- phylogeny
- herbivory

THANK YOU FOR ATTENTION.

**Questions
and/or
comments?**