

# Summary

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# Introduction

Functional Ecology aims to study organisms in terms of functional traits [citation needed], instead of studying species.

From Hutchinson definition of *fundamental niche* (Hutchinson, 1957), Violle and Jiang (2009) extended the definition to become a *functional niche*—a multi-dimensional volume, also called the *functional space* ; it is an  $n$ -dimensional space defined by measures on  $n$  traits. In such a space an individual is defined by all the values of its traits, each one on a distinct axis. In three dimensions, the species niche would be the volume encompassing all its individuals; the center of gravity of the cloud would be the species average trait values, defining niche position, while the shape of the general "cloud" would define niche breadth, as suggested by Violle and Jiang (2009).

The concept of functional niche helps understand community ecology and assemblage. In this view, two species would coexist if their functional volumes would not intersect. The high number of dimensions, i.e. traits, of the functional "hyper-volume" as called by [citation needed] makes it difficult to apprehend: the volume may have "holes" where some trait combinations are impossible. An individual is located by its own trait values or relatively to its species average. The distance from species average translates the intra-specific variability in the species.

Species functional niche reflects their ecological strategies. For plants, four traits have been identified to underline distinct strategies: the classical Leaf Area - Height - Seed mass triangle, suggested by [citation needed]; as well as wood economics spectrum traits (Baraloto et al., 2010a).

Functional trait → performance indexes → fitness, performance indexes reflect fitness, thus evolution. Linking functional traits and performance indexes → predict the evolution of functional traits. How then species position do not change over time to increase performance? Some case, such as wood density, where we would expect species to decrease it to have faster growth two answers: resource-energy trade-offs (darwinian demons) or specific strategy.

Those trade-offs are generally shown at the species level by comparing several species showing impossible trait combinations, i.e. in the previous example it would be impossible to find species with dense wood and fast growth. However, if we want to investigate not a general trade-off that exists at the species level, but how the position of an individual compared to the species average may influence performance indexes. For example, a tree with a less dense wood than its species average would have a higher growth rate than individuals with a density around the species average. Throughout this paper we try to understand how the performance of an individual is affected by the distance to its species average trait.

For certain traits, we would expect the species position not to vary, thus, our approach may unravel new trade-offs between intra-specific and inter-specific variabilities.

Diameter Growth & Tropical Forests → French Guiana Context

Several growth models created, used → estimate growth using measured traits

# Materials and Methods

## Data Provenance

### Growth Data

The first data set is an inventory of all trees over 10cm in Diameter at Breast Height (DBH), i.e. measured at 1.3m high, in nine 1-ha plots in French Guiana (see map [\[missing figure\]](#)). In each plot, trees diameter were measured every two or five years depending on the plot.

We selected a common measured period between 2001 and 2013 comprising a total of 3549 trees; we estimated annual growth rate (AGR) in diameter by fitting a linear regression of DBH over years. The slope of the regression gave us an "average" AGR for each followed tree on the comprised

### Trait Data

The second data set comes was a collection of five functional traits (see [Table 1](#)) extracted from a bigger database ([Baraloto et al., 2010a,b](#)) on the same trees. Traits were not followed through time and measured only once. Selected traits are related to leaf and wood economics spectrum.

## Growth model

To predict the growth of tree from a single trait we used a linear mixed-model with the general formula:

$$\log(\text{AGR}_{i,s,p} + 1) = \theta_0 + \gamma_{0,s} + \gamma_p + (\theta_1 + \gamma_{1,s}) \times \text{DBH} + (\theta_2 + \gamma_{2,s}) \times \log(\text{DBH}) + \delta + \epsilon_i, \quad (1)$$

with  $\epsilon_i \sim \mathcal{N}(0, \theta_3)$  the individual residual, where  $\text{AGR}_{i,s,p}$  is the AGR of tree  $i$  of species  $s$  in plot  $p$ ;  $\theta_0 \dots \theta_3$  are parameters to be estimated;  $\gamma_{0,s} \dots \gamma_{2,s}$  and  $\gamma_p$  follow a centered Gaussian distribution with unknown variances  $\sigma_{0,s}^2 \dots \sigma_{2,s}^2$  and  $\sigma_p^2$ ;  $\text{Tr}_s$  is the average trait value for species  $s$ .

To understand how the distance to species average value affected the predicted growth, we used different  $\delta$  values:  $\theta_4 \times \text{Tr}_s$  the species average value, with a parameter  $\theta_4$ ;  $\theta_4 \times \text{Tr}_s + \theta_5 \times (\text{Tr}_i - \text{Tr}_s)$  the distance of individual trait value  $\text{Tr}_i$  to species average, with the species term  $\text{Tr}_s$ ; or  $\theta_4 \times \text{Tr}_s + \theta'_5 \times |\text{Tr}_i - \text{Tr}_s|$  the absolute distance to species average trait, with the species term  $\text{Tr}_s$ ; or  $\theta'_4 \times \text{Tr}_i$  the individual trait value.

Our models tested the difference of prediction between using only the species average trait value to predict growth and the same term plus an individual distance term (real or absolute) vs. the individual trait value.

## Data analysis

All data analyses were made using [R Core Team \(2015\)](#) version 3.2.0 (2015-04-16), plots were made with [Wickham \(2009\)](#). We fit mixed-models with "lme4" R package ([Bates et al., 2014](#)) 1.1-7 and computed adapted R-squared for mixed-models ([Nakagawa and Schielzeth, 2013](#)) implemented in "MuMIn" R package ([Bartón, 2015](#)) version 1.13.4.

Trait Name	Units	Role
Trunk bark thickness	mm	Defence, Stem economics spectrum
Xylem density (wood density)	$\text{g.cm}^{-3}$	Stem economics
Specific Leaf Area (SLA)	$\text{cm}^2.\text{g}^{-1}$	Leaf economics
Laminar total chlorophyll	$\mu\text{m}.\text{mm}^{-2}$	Leaf economics
Laminar toughness	N	Leaf economics

**Table 1:** Selected functional traits.

**Results**

**Discussion**

**Authors Contributions and Acknowledgments**

## References

- Baraloto, C., Timothy Paine, C. E., Patiño, S., Bonal, D., Hérault, B. and Chave, J. (2010a). Functional trait variation and sampling strategies in species-rich plant communities. *Functional Ecology* 24, 208--216.
- Baraloto, C., Timothy Paine, C. E., Poorter, L., Beauchene, J., Bonal, D., Domenach, A.-M., Hérault, B., Patiño, S., Roggy, J.-C. and Chave, J. (2010b). Decoupled leaf and stem economics in rain forest trees. *Ecology Letters* 13, 1338--1347.
- Bartoń, K. (2015). MuMIn: Multi-Model Inference. R package version 1.13.4.
- Bates, D., Mächler, M., Bolker, B. and Walker, S. (2014). Fitting Linear Mixed-Effects Models using lme4. *arXiv:1406.5823 [stat]* .
- Hutchinson, G. E. (1957). Concluding Remarks. *Cold Spring Harb Symp Quant Biol* 22, 415--427.
- Nakagawa, S. and Schielzeth, H. (2013). A general and simple method for obtaining R<sup>2</sup> from generalized linear mixed-effects models. *Methods Ecol Evol* 4, 133--142.
- R Core Team (2015). R: A Language and Environment for Statistical Computing. R Foundation for Statistical Computing Vienna, Austria.
- Violle, C. and Jiang, L. (2009). Towards a trait-based quantification of species niche. *J Plant Ecol* 2, 87--93.
- Wickham, H. (2009). *ggplot2: elegant graphics for data analysis*. Springer New York.