SUPPLEMTARY MATERIAL

Riofrío, J. *et al.* (2019) Species mixing effects on height-diameter and basal area increment models for Scots pine and Maritime pine. *Forests*.



Figure S1. Tree height plotted against diameter of Scots pine (*Pinus sylvestris*) and Maritime pine (*Pinus pinaster*) in mixed (black open symbols) and pure stands (gray open symbols) for the triplets network dataset (Training dataset).



Figure S2. Basal area increment data used for models fitting in 5 cm diameter classes for Scots pine (*Pinus sylvestris*) and Maritime pine (*Pinus pinaster*) in mixed and pure stands.



Figure S3. Tree height plotted against diameter of *Pinus sylvestris* and *Pinus pinaster* in mixed (black open symbols) and pure stands (gray open symbols) from National Forest Inventory data (Validation dataset).



Figure S4. Plot-specific *h*-*d* curves using fitted models overlaid on the measured tree heights (gray dots) for Scots pine and Maritime pine.



Figure. S5. Standardized residuals of generalized non-linear mixed-effects *h*-*d* models for Scots pine and Maritime pine.



Figure S6. Mean 5-year basal area increment of *Pinus sylvestris* and *Pinus pinaster* in mixed and pure stands.



Figure S7. Effects of stand basal area (BA), basal area of larger trees (BAL) and crown ratio on the final basal area increment model for Scots pine (Table 4). Curves were produced using the parameter estimates in Table 2 and varying one explanatory variable at a time. Mean values of the data were used for predictors and the range of the variable of interest in the figure.



Figure S8. Differences of dominant height of each species between mixed and pure stands. Mean and standard error for Scots pine (white circles) and Maritime pine (black circles).

Appendix S1.

Screening for generalized h-d models

In this study we used nonlinear functions widely used for modelling *h*-*d* relationships. We evaluated functions tested previously to fit the *h*-*d* relationships for both species in mono-specific stands (M1-M5 in Table S1) and asymptotic general functions used for other species (M6-M9 in Table S1). The basic function forms were expanded for including covariates that described stand characteristics, which would have significant influences on the *h*-*d* relationships and improve tree height estimates. This was accomplished by expressing the asymptotic parameter in each equation (α_0 parameter) as a function of the selected covariates.

Function	in Function form Parameter expansion		Source base
number			function
M1	$h = 13 + \alpha_0 \ e^{\left(\frac{-\alpha_1}{\sqrt{a}}\right)}$	$\alpha_0 = \beta_0 + \beta_1 H o + \beta_2 V_2 + \ldots + \beta_i V_i$	Schöder & Álvarez (2001)
M2	$h = \alpha_0 \; e^{\left(\frac{\alpha_1}{d}\right)}$	$\alpha_0 = \beta_0 H o^{\beta_1} \cdot V_2^{\beta_2} \cdot \dots \cdot V_i^{\beta_i}$	del Río (1999)
M3	$h = \alpha_0 \ e^{\left[\alpha_1 \left(\frac{1}{d} - \frac{1}{Do}\right)\right]}$	$\alpha_0 = \beta_0 H o^{\beta_1} \cdot V_2^{\beta_2} \cdot \dots \cdot V_i^{\beta_i}$	del Río (1999)
M4	$h = Ho \ e^{\left[\alpha_0 \cdot \left(\frac{1}{d} - \frac{1}{Do}\right)\right]}$	$\alpha_0 = \beta_1 H o + \beta_2 V_2 + \ldots + \beta_i V_i$	Michailoff (1943)
M5	$h = Ho \ e^{\left[\alpha_0 \cdot \left(\frac{1}{d} - \frac{1}{Do}\right)\right]}$	$\alpha_0 = \beta_0 + \beta_1 H o + \beta_2 V_2 + \ldots + \beta_i V_i$	Michailoff (1943)
M6	$h = 1.3 + \alpha_0 \left[1 - e^{(\alpha_1 \cdot d)} \right]^{\alpha_2}$	$\alpha_0 = \beta_0 + \beta_1 H o + \beta_2 V_2 + \ldots + \beta_i V_i$	Richards (1959)
M7	$h=1.3+\alpha_0\left[1-e^{(\alpha_1\cdot d^{\alpha_2})}\right]$	$\alpha_0 = \beta_0 + \beta_1 H o + \beta_2 V_2 + \ldots + \beta_i V_i$	Yang (1978)
M8	$h = 1.3 + \alpha_0 e^{(\alpha_1 \cdot d^{-\alpha_2})}$	$\alpha_0 = \beta_0 + \beta_1 H o + \beta_2 V_2 + \ldots + \beta_i V_i$	Korf (Zeide 1989)
M9	$h = 1.3 + e^{\left(\alpha_0 + \frac{\alpha_1}{d + \alpha_2}\right)}$	$\alpha_0 = \beta_0 + \beta_1 H o + \beta_2 V_2 + \ldots + \beta_i V_i$	Ratkowsky (1990)

Table S1. Generalized height-diameter equations evaluated.

h: tree height (m); *d*: diameter at breast height (cm); α_0 , α_1 , α_2 are species dependent parameters. β_0 and $\beta_{1....i}$ are species model parameters for each explanatory variables tested in the model selection procedure. Note that Table S1 contains only the basic fixed effects models of the *h*-*d* analyzed in this study.

References

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Río, M. del. 1999. Régimen de claras y modelo de producción para *Pinus sylvestris* L. en los Sistemas Central e Ibérico. Tesis Doctorales INIA nº2. Serie Forestal. 257 pp

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Zeide B (1989) Accuracy of equations describing diameter growth. Can J For Res 19:1283–1286

Maritime pine			Scots pine		
Model	AIC	RMSE	Model	AIC	RMSE
M3	2880.4	1.362	M3	3388.2	1.568
M2	2890.3	1.380	M2	3390.1	1.569
M4	2895.3	1.377	M7	3390.4	1.571
M9	2896.5	1.379	M9	3392.7	1.584
M5	2897.8	1.376	M6	3393.2	1.571
M7	2907.1	1.376	M4	3396.9	1.573
M8	2914.5	1.379	M5	3397.5	1.573
M6	2914.8	1.381	M8	3400.7	1.576
M1	2925.2	1.392	M1	3401.2	1.576

Table S2. Ranking and summary statistics for generalized height–diameter models fitted without considering the species-mixing effects, according to functions of Table S1

Appendix S2.

Models selection

To evaluate the performance among different set of *h*-*d* functions and individual tree growth models, we used the multi-model inference approach based on information-theory (Anderson, 2007), which allows evaluation of multiple non-nested models relative to each other and quantification of the relative support for multiple models simultaneously. Fitted models were ranked by lowest AICc (Second-order Akaike Information Criterion) and greater Akaike weights (w_i) values. The following Akaike weight of each model is a measure of the probability that model *i* is the best model, e.g., the most parsimonious, for the observed data and given the set of candidate models:

$$w_{i} = \frac{exp\left\{-\frac{1}{2}\Delta_{i}(AICc)\right\}}{\sum_{k=1}^{K}exp\left\{-\frac{1}{2}\Delta_{k}(AICc)\right\}}$$
(1)

where Δ is the difference in AICc between model *i* and the best candidate model for *i* = 1, ... *K*. Therefore, the final output is a set of candidate models rather than a single model.

References

Anderson, D. R. Model Based Inferences in the Life Sciences: a Primer on Evidence; Springer, 2007; ISBN 9780387740737.