

Appendices: Lianas affect population growth rates of tropical trees

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S1 TEXT

S1.1 DETAILS ON VITAL RATE ESTIMATION

Probability of reproduction: We estimated the size-dependent probability of reproduction (R) using a GLMM, where tree reproductive status (sterile or reproductive; datasets 2 and 3) was related (logistically) to tree diameter and level of liana infestation (eq. 1). Reproductive status was dichotomized for the analyses using only sterile trees (score 0) versus fertile trees with reproductive structures (score 1-4).

Fraction of crown bearing fruits: We used a linear mixed effect model to examine the relationship between field assessments of the fraction of the crown bearing reproductive structures (hereafter crown reproductive fraction; C) and tree diameter and the level of liana infestation (eq. 1; datasets 2 and 3).

Seed production: Mean species-specific seed production per m² of reproductive basal area (F) was quantified by dividing seed production by reproductive basal area. Reproductive basal area was calculated for each census year as the sum of individual basal areas (dataset 1) weighted by size-dependent reproductive probability and crown reproductive fraction (datasets 2 and 3) and interpolated linearly between census years to obtain yearly values. Yearly seed production equaled plot area multiplied by the density of seeds arriving in the seed traps (dataset 4). Yearly estimates of seed production were divided by yearly estimates of reproductive basal area and averaged to obtain F.

Seedling establishment: We calculated species-specific mean yearly seedling establishment probability (ϕ) as the density (per area) of newly recruiting seedlings across all seedling plots divided by the density of seeds arriving at all seed traps (datasets 4 and 5).

Initial height of establishing seedlings: The initial height distributions of establishing tree seedlings (I; dataset 5) were fit to exponential, log-normal and Weibull probability density functions. The Weibull provided the best fit for 31 out of 33 species. We therefore fit Weibull distributions for every species.

Seedling to tree transition: We estimated the height at which tree seedlings entered the FDP census (hd=10; 10 mm DBH) with a linear model that related height with DBH for each species using data on height and DBH from seedling data (dataset 6).

Growth: We modeled growth (G) as height growth for seedlings (mm/year) and basal area growth for trees > 1 cm DBH (mm^2 /year) using linear mixed effect models (eq. 1). We calculated growth rates as the difference in size divided by the time in years between censuses (datasets 1, 5 and 6). Due to

measurement error (Ruger et al. 2011), a tiny fraction of the growth rates were unrealistically low or high (dataset 1), and likely erroneous, we therefore excluded any values greater than four standard deviations from the overall mean (0.4% of all measurements).

Survival: We estimated the size-dependent yearly probability of survival (S) for both seedlings and larger individuals (> 1 cm DBH) using logistic mixed effects models (datasets 1, 5 and 6). Tree survival rates were related to L as shown in eq. 1.

Minimum and maximum sizes: Minimum sizes (A) in the IPM were set to 0 mm height for seedlings and 10 mm DBH for larger trees. Maximum sizes (Z) were the previously estimated species-specific height at which seedlings have 10 mm DBH for seedlings and 110% (sensu Zuidema et al. 2010) of the mean DBH of the 6 largest individuals of each species in the 50-ha plot (dataset 1).

S2 TABLES

Table S1. Overview of species used in each analysis, ordered alphabetically. A cross (X) indicates a species was included.

Species	Fecundity	Reproduction	Growth	Survival
<i>Acalypha diversifolia</i>	X	X		
<i>Alchornea costaricensi</i>	X	X	X	X
<i>Alseis blackiana</i>	X	X	X	X
<i>Anacardium excelsum</i>	X	X	X	X
<i>Apeiba membranacea</i>	X	X	X	X
<i>Apeiba tibourbou</i>	X	X	X	X
<i>Aspidosperma spruceanum</i>	X	X		
<i>Astrocaryum standleyanum</i>	X	X	X	X
<i>Astronium graveolens</i>	X	X	X	X
<i>Beilschmiedia pendula</i>	X	X	X	X
<i>Brosimum alicastrum</i>	X	X		
<i>Calophyllum longifolium</i>	X	X		
<i>Casearia aculeata</i>	X	X		
<i>Casearia arborea</i>	X	X	X	X
<i>Cassipourea elliptica</i>	X	X		
<i>Cecropia insignis</i>	X	X		
<i>Cecropia obtusifolia</i>	X	X	X	X
<i>Ceiba pentandra</i>	X	X	X	X
<i>Chrysophyllum argenteum</i>	X	X	X	X
<i>Chrysophyllum cainito</i>	X	X	X	X
<i>Cordia alliodora</i>	X	X		
<i>Cordia bicolor</i>	X	X		
<i>Coussarea curvigemma</i>	X	X		
<i>Croton billbergianu</i>	X	X		
<i>Cupania seemannii</i>	X	X	X	X
<i>Desmopsis panamensis</i>	X	X	X	X
<i>Dipteryx oleifera</i>	X	X		
<i>Drypetes standleyi</i>	X	X		

Continued on next page

S2 TABLES

Table S1. Overview of species used in each analysis, ordered alphabetically. A cross (X) indicates a species was included.

Species	Fecundity	Reproduction	Growth	Survival
<i>Eugenia galalonensis</i>	X	X	X	X
<i>Eugenia nesiotica</i>	X	X		
<i>Eugenia oerstediana</i>	X	X	X	X
<i>Faramea occidentalis</i>	X	X	X	X
<i>Garcinia intermedia</i>	X	X	X	X
<i>Guapira standleyana</i>	X	X		
<i>Guarea guidonia</i>	X	X	X	X
<i>Guatteria dumetorum</i>	X	X		
<i>Guettarda foliacea</i>	X	X	X	X
<i>Gustavia superba</i>	X	X	X	X
<i>Hasseltia floribunda</i>	X	X	X	X
<i>Heisteria concinna</i>	X	X		
<i>Hirtella triandra</i>	X	X	X	X
<i>Hura crepitans</i>	X	X		
<i>Hieronyma alchorneoides</i>	X	X	X	X
<i>Inga marginata</i>	X	X	X	X
<i>Jacaranda copaia</i>	X	X	X	X
<i>Lactia thamnina</i>	X	X		
<i>Lindackeria laurina</i>	X	X		
<i>Lonchocarpus heptaphyllus</i>	X	X	X	X
<i>Luehea seemannii</i>	X	X		
<i>Macrocnemum roseum</i>	X	X		
<i>Mosannonia garwoodii</i>	X	X		
<i>Miconia affinis</i>	X	X	X	X
<i>Miconia argentea</i>	X	X	X	X
<i>Miconia nervosa</i>	X	X		
<i>Mouriri myrtilloides</i>	X	X	X	X
<i>Nectandra cissiflora</i>	X	X		
<i>Ocotea cernua</i>	X	X	X	X
<i>Ocotea oblonga</i>	X	X	X	X
<i>Ocotea whitei</i>	X	X	X	X
<i>Oenocarpus mapora</i>	X	X		
<i>Platymiscium pinnatum</i>	X	X		
<i>Platydictyon elegans</i>	X	X	X	X
<i>Pachira sessilis</i>	X	X	X	X
<i>Poulsenia armata</i>	X	X		
<i>Pouteria reticulata</i>	X	X	X	X
<i>Prioria copaifera</i>	X	X		
<i>Protium costaricense</i>	X	X		
<i>Protium panamense</i>	X	X	X	X
<i>Protium tenuifolium</i>	X	X	X	X
<i>Pseudobombax septenatum</i>	X	X		
<i>Pterocarpus rohrii</i>	X	X	X	X
<i>Quararibea asterolepis</i>	X	X	X	X
<i>Randia armata</i>	X	X	X	X
<i>Rinorea sylvatica</i>	X	X	X	X
<i>Attalea butyracea</i>	X	X		
<i>Simarouba amara</i>	X	X		
<i>Sloanea terniflora</i>	X	X	X	X
<i>Socratea exorrhiza</i>	X	X	X	X
<i>Spondias radlkoferi</i>	X	X	X	X
<i>Tabebuia guayacan</i>	X	X	X	X

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Table S1. Overview of species used in each analysis, ordered alphabetically. A cross (X) indicates a species was included.

Species	Fecundity	Reproduction	Growth	Survival
<i>Tabebuia rosea</i>	X	X	X	X
<i>Tabernaemont arborea</i>	X	X	X	X
<i>Tachigali versicolor</i>	X	X	X	X
<i>Talisia nervosa</i>	X	X		
<i>Terminalia amazonia</i>	X	X	X	X
<i>Terminalia oblonga</i>	X	X		
<i>Tetragastris panamensis</i>	X	X		
<i>Trattinnicki aspera</i>	X	X	X	X
<i>Trichilia pallida</i>	X	X		
<i>Trichilia tuberculata</i>			X	X
<i>Triplaris cumingiana</i>			X	X
<i>Unonopsis pittieri</i>			X	X
<i>Viola sebifera</i>			X	X
<i>Viola multiflora</i>			X	X
<i>Viola surinamensis</i>			X	X
<i>Vochysia ferruginea</i>			X	X
<i>Xylopia macrantha</i>			X	X
<i>Zanthoxylum ekmanii</i>			X	X
<i>Zanthoxylum panamense</i>			X	X

Table S2. The focal species used to create integral projection models, and corresponding sample sizes in the various datasets (described in the maintext). The columns seeds, recruits, seedlings, trees, and reproduction correspond to the number of individual records of seeds (dataset 2), new recruits (dataset 3) from the seedling census (dataset 4), BCI 50ha FDP census and towerplot data (dataset 1), and the data on the reproduction and liana census (dataset 6). A total of 33 have enough data

Species	Seeds	Recruits	Seedlings	Trees	Reproduction
<i>Alchornea costaricensi</i>	10187	77	336	828	215
<i>Alseis blackiana</i>	152070	70	2060	13182	1203
<i>Apeiba membranacea</i>	8028	127	298	570	160
<i>Beilschmiedia pendula</i>	1708	1935	16382	4056	77
<i>Brosimum alicastrum</i>	6687	97	1558	1194	262
<i>Cecropia insignis</i>	80899	187	834	2551	684
<i>Chrysophyllum argenteum</i>	39	33	407	1119	45
<i>Chrysophyllum cainito</i>	919	220	908	271	22
<i>Cordia alliodora</i>	5188	105	477	433	227
<i>Cordia bicolor</i>	5770	63	641	2222	503
<i>Drypetes standleyi</i>	639	81	1122	3080	129
<i>Eugenia oerstediana</i>	1717	883	6906	4039	91
<i>Garcinia intermedia</i>	1237	53	971	6945	792
<i>Guarea guidonia</i>	216	365	1013	3134	424
<i>Guatteria dumetorum</i>	907	505	191	2293	167
<i>Gustavia superba</i>	741	222	1589	1187	57
<i>Heisteria concinna</i>	79938	160	1072	1189	219
<i>Hieronyma alchorneoides</i>	155	104	70	265	36
<i>Jacaranda copaia</i>	97	43	284	783	416
<i>Luehea seemannii</i>	3385	115	540	648	508
<i>Miconia argentea</i>	117	186	728	2884	177
<i>Ocotea whitei</i>	350	68	4715	1566	261
<i>Platypodium elegans</i>	646	340	105	312	132
<i>Pouteria reticulata</i>	352	111	3869	2705	190
<i>Quararibea asterolepis</i>	1213	184	7317	3207	52
<i>Simarouba amara</i>	1472	98	784	3787	381
<i>Tabebuia guayacan</i>	3262	133	254	132	132
<i>Tabebuia rosea</i>	730	44	464	573	279
<i>Tabernaemontana arborea</i>	3729	58	365	2544	78
<i>Tetragastris panamensis</i>	2343	575	3909	7213	327
<i>Trichilia pallida</i>	366	49	146	1157	500
<i>Trichilia tuberculata</i>	28841	3756	9577	20309	508
<i>Triplaris cumingiana</i>	716	154	247	640	355

Table S3. Sample size within each liana infestation class (see main document), for seed production and reproductive fraction, for each of focal species used to create integral projection models. The columns L0-L4 correspond to 0%, 1% - 24%, 25% - 49%, 49% - 74% and 75% - 100% of the crown area infested with lianas.

Species	L0	L1	L2	L3	L4
<i>Alchornea costaricensi</i>	37	47	47	27	53
<i>Alseis blackiana</i>	217	162	117	77	212
<i>Apeiba membranacea</i>	53	46	27	18	27
<i>Beilschmiedia pendula</i>	27	22	6	6	16
<i>Brosimum alicastrum</i>	116	69	34	29	34
<i>Cecropia insignis</i>	576	12	2	0	11
<i>Chrysophyllum argenteum</i>	5	8	2	3	24
<i>Chrysophyllum cainito</i>	3	6	5	4	2
<i>Cordia alliodora</i>	116	22	6	3	12
<i>Cordia bicolor</i>	169	77	99	80	75
<i>Drypetes standleyi</i>	64	34	13	10	8
<i>Eugenia oerstediana</i>	36	15	16	17	7
<i>Garcinia intermedia</i>	408	208	90	37	83
<i>Guarea guidonia</i>	304	177	34	27	27
<i>Guatteria dumetorum</i>	69	34	28	23	33
<i>Gustavia superba</i>	26	18	6	4	3
<i>Heisteria concinna</i>	109	120	48	18	19
<i>Hieronyma alchorneoides</i>	271	235	63	55	48
<i>Jacaranda copaia</i>	243	28	13	11	20
<i>Luehea seemannii</i>	72	113	68	52	65
<i>Miconia argentea</i>	56	27	25	9	60
<i>Ocotea whitei</i>	88	74	49	31	19
<i>Platypodium elegans</i>	29	20	13	8	27
<i>Pouteria reticulata</i>	78	40	18	15	32
<i>Quararibea asterolepis</i>	28	8	6	3	4
<i>Simarouba amara</i>	176	43	31	40	80
<i>Tabebuia guayacan</i>	31	14	14	10	15
<i>Tabebuia rosea</i>	64	31	15	20	86
<i>Tabernaemontana arborea</i>	22	26	10	8	12
<i>Tetragastris panamensis</i>	99	112	70	46	48
<i>Trichilia pallida</i>	197	152	77	38	36
<i>Trichilia tuberculata</i>	229	125	89	53	36
<i>Triplaris cumingiana</i>	277	11	19	6	19

Table S4. Sample size within each liana infestation class (see main document), for growth and survival, for each of focal species used to create integral projection models. The columns L0-L4 correspond to 0%, 1% - 24%, 25% - 49%, 49% - 74% and 75% - 100% of the crown area infested with lianas.

Species	L0	L1	L2	L3	L4
<i>Alchornea costaricensi</i>	14	39	47	18	29
<i>Alseis blackiana</i>	52	54	34	32	99
<i>Apeiba membranacea</i>	4	1	6	8	23
<i>Beilschmiedi pendula</i>	8	3	5	2	11
<i>Brosimum alicastrum</i>	44	10	10	14	12
<i>Cecropia insignis</i>	189	6	2	1	2
<i>Chrysophyllu argenteum</i>	3	12	13	4	23
<i>Chrysophyllu cainito</i>	3	5	6	4	2
<i>Cordia alliodora</i>	69	10	8	3	9
<i>Cordia bicolor</i>	98	54	79	63	75
<i>Drypetes standleyi</i>	4	4	7	4	6
<i>Eugenia oerstediana</i>	41	18	18	17	11
<i>Garcinia intermedia</i>	162	99	68	26	45
<i>Guarea guidonia</i>	8	9	19	7	15
<i>Guatteria dumetorum</i>	12	5	4	1	4
<i>Gustavia superba</i>	50	44	20	14	29
<i>Heisteria concinna</i>	6	15	2	4	8
<i>Hieronyma alchorneoide</i>	7	13	6	9	11
<i>Jacaranda copaia</i>	185	20	14	7	14
<i>Luehea seemannii</i>	28	32	31	23	41
<i>Miconia argentea</i>	60	28	26	9	64
<i>Ocotea whitei</i>	17	30	28	19	13
<i>Platypodium elegans</i>	6	11	2	1	13
<i>Pouteria reticulata</i>	4	7	5	2	8
<i>Quararibea asterolepis</i>	16	14	9	5	10
<i>Simarouba amara</i>	103	24	20	21	38
<i>Tabebuia guayacan</i>	11	5	10	4	6
<i>Tabebuia rosea</i>	13	13	7	4	34
<i>Tabernaemont arborea</i>	5	8	6	6	17
<i>Tetragastris panamensis</i>	8	23	25	20	41
<i>Trichilia pallida</i>	73	81	48	23	11
<i>Trichilia tuberculata</i>	28	40	29	21	64
<i>Triplaris cumingiana</i>	117	6	10	1	11

Table S5. AIC values for the size dependent fraction of reproductive individuals, including Δ AIC values. Model null, size, main and full correspond to mixed effect models including no predictions, using only size, size including additive effects of liana load and full interactions between size and liana load respectively.

Model	AIC	Δ AIC
Null	19219	3512
Size	16498	791
Main	15710	3
Full	15707	0

Table S6. AIC values for the size dependent seed production (fraction of crown bearing reproductive structures), including Δ AIC values. Model null, size, main and full correspond to mixed effect models including no predictions, using only size, size including additive effects of liana load and full interactions between size and liana load respectively.

Model	AIC	Δ AIC
Null	2300	450
Size	2082	233
Main	1850	0
Full	1875	25

Table S7. AIC values for the size dependent basal area growth, including Δ AIC values. Model null, size, main and full correspond to mixed effect models including no predictions, using only size, size including additive effects of liana load and full interactions between size and liana load respectively.

Model	AIC	Δ AIC
Null	94670	11665
Size	94543	11538
Main	83062	57
Full	83005	0

Table S8. AIC values for the size dependent survival, including Δ AIC values. Model null, size, main and full correspond to mixed effect models including no predictions, using only size, size including additive effects of liana load and full interactions between size and liana load respectively.

Model	AIC	Δ AIC
Null	5191	587
Size	5075	471
Main	4603	0
Full	4609	6

Table S9. Fit coefficients (fixed-effects) of the AIC selected mixed models (and standard errors). Intercept and slope values are the "species mean" coefficients relating to size-dependent reproduction, fecundity, growth and survival for liana free trees. Coefficients L1-L4 relate to main effects of lianas for each crown infestation class (see main text), while the L(1-4):size are the coefficients relating to size - infestation class interactions.

	Fecundity	Reproduction	Growth	Survival
Intercept	0.6833 (0.0224)	-0.4628 (0.2924)	2772.214 (308.1966)	2.067 (0.1346)
Slope	0.1762 (0.0218)	2.8014 (0.2603)	-37.3389 (193.466)	-0.1963 (0.1003)
L1	-0.0763 (0.0132)	0.034 (0.0909)	-443.3472 (135.146)	0.1746 (0.1549)
L2	-0.1141 (0.0215)	-0.2458 (0.1182)	-855.2915 (199.6878)	0.0702 (0.1717)
L3	-0.167 (0.0233)	-0.5911 (0.1321)	-999.2213 (179.128)	-0.2263 (0.1926)
L4	-0.1895 (0.0338)	-1.9947 (0.1899)	-1337.3534 (200.4283)	-0.5799 (0.179)
L1:size			103.2487 (127.7744)	
L2:size			395.7886 (159.3142)	
L3:size			-529.8031 (175.0574)	
L4:size			-207.356 (160.3242)	

S3 FIGURES

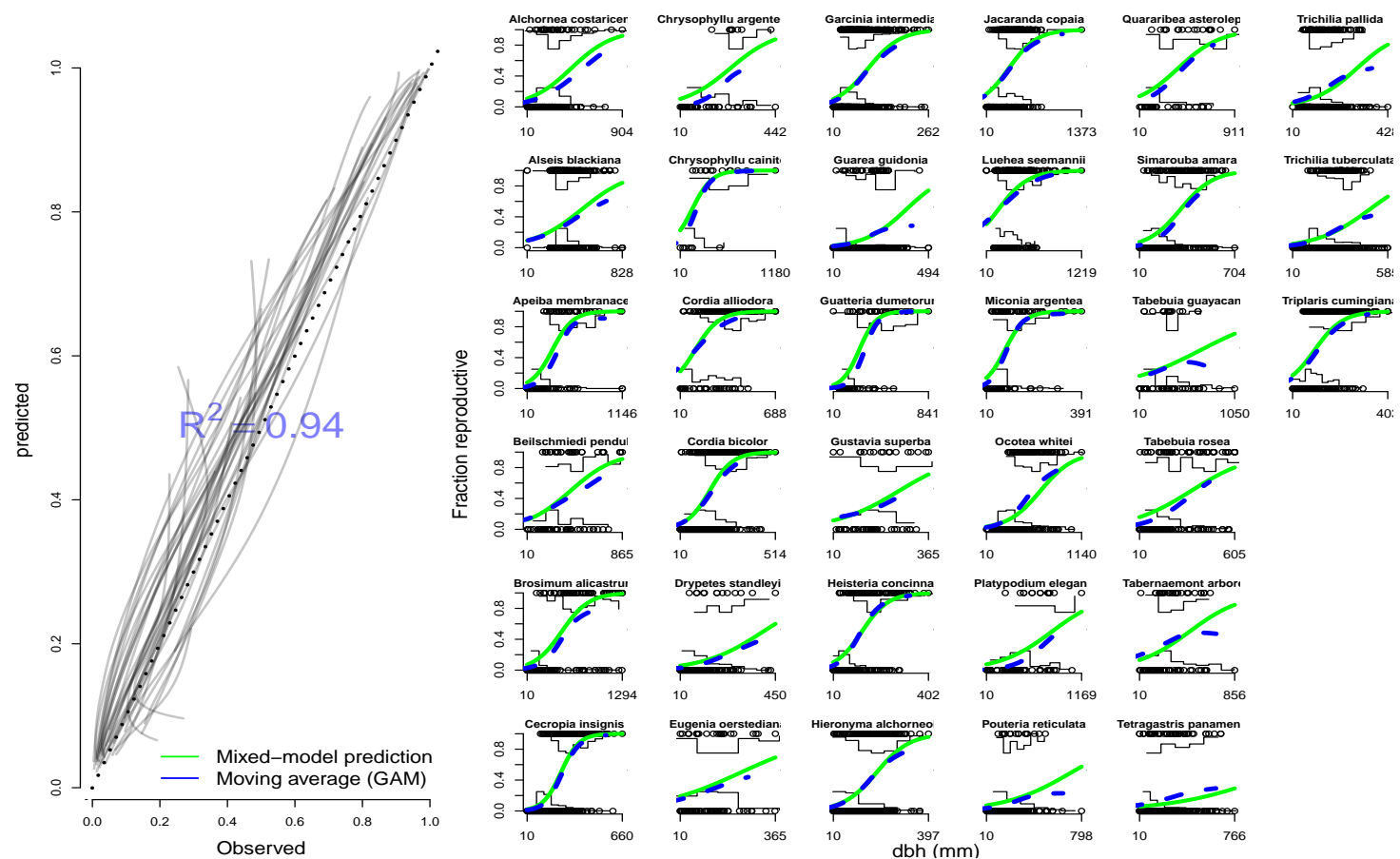


Figure S1. The proportion of reproductive individuals as a function of tree size (dbh). The large panel on the left shows the agreement between the model predictions (y-axis) from the mixed-model (see maintext) and an general additive model (GAM) fit for each species on the data (x-axis). The smaller panels on the right show model fits per species over size (dbh in mm). Here, the green line shows the predicted values by the fitted mixed effect model (corresponding to the y-axis on the large left panel), while the blue lines indicates a GAM predicted moving average (corresponding to the x-axis on the large left panel). The step-wise lines are indicative of the relative density of the reproductive status at a given size.

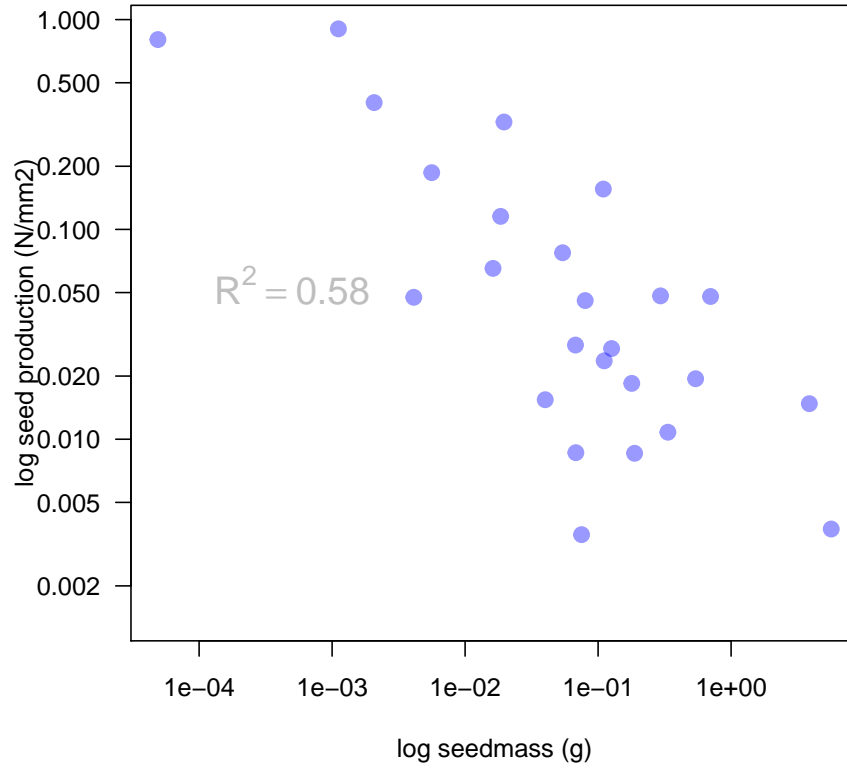


Figure S2. Estimates of seed production, in numbers of seeds produced per unit basal area plotted against seed dry mass (g) from Wright *et al.* 2010.

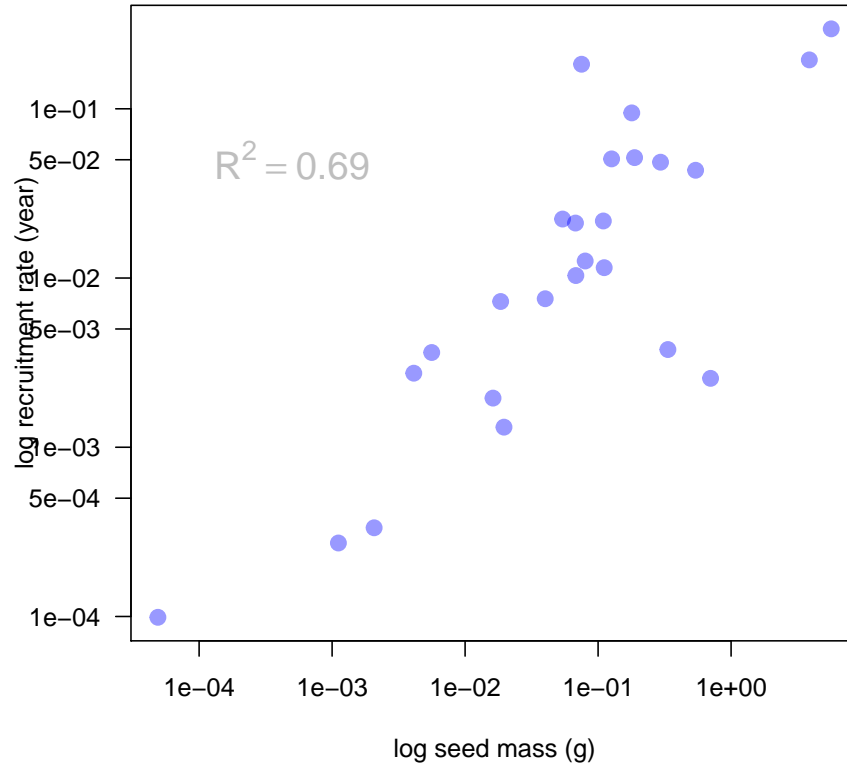


Figure S3. Estimates of the annual seed to seedling transition rate plotted against seed dry mass (g) from Wright *et al.* 2010.

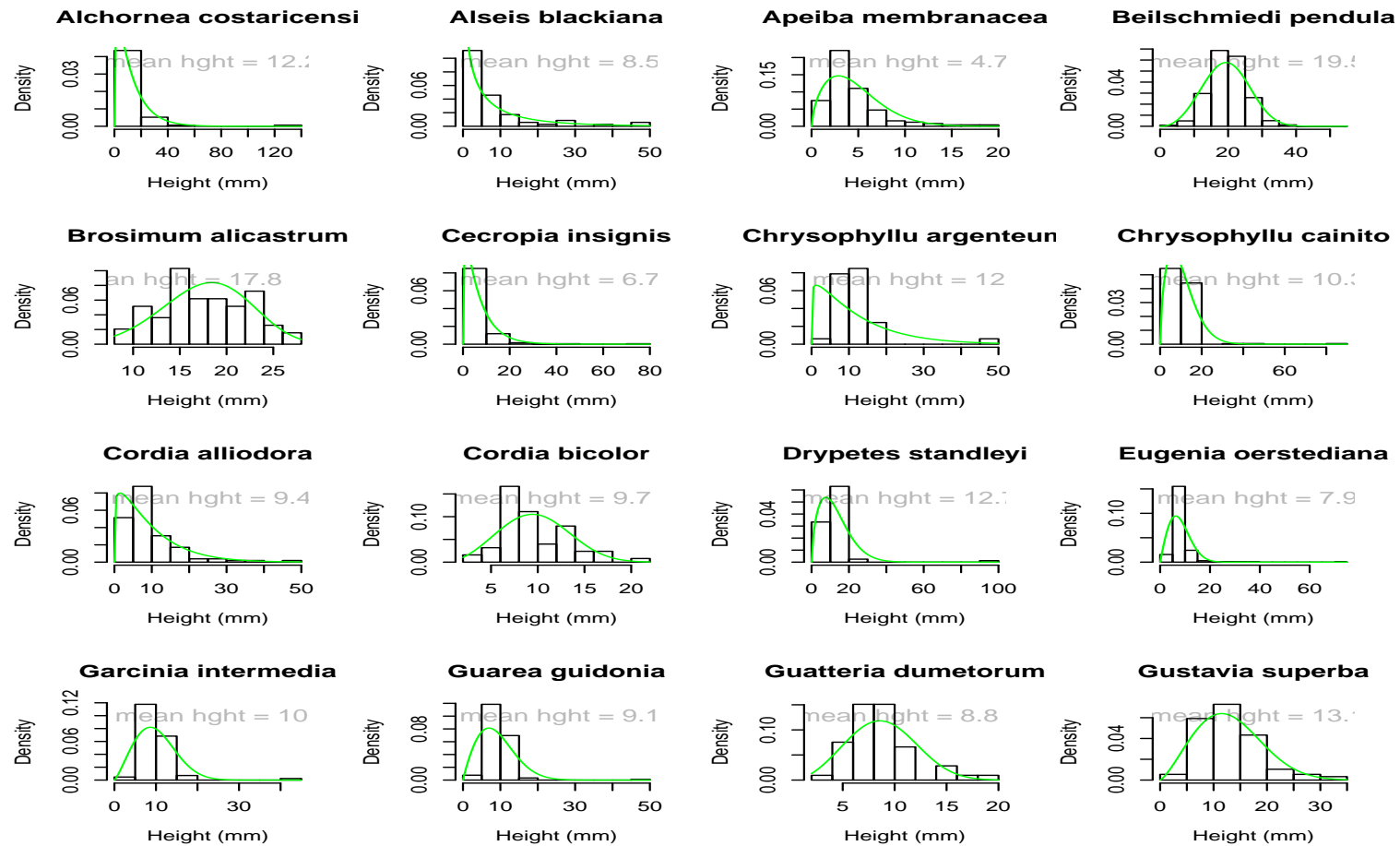


Figure S4. Distribution of initial heights of establishing seedlings, including fit models (green lines). Seedling heights of zero indicate very small seedlings that could not be measured without danger of inflicting damage. The above multi-panel graph is number 1 of 3 plots containing all species.

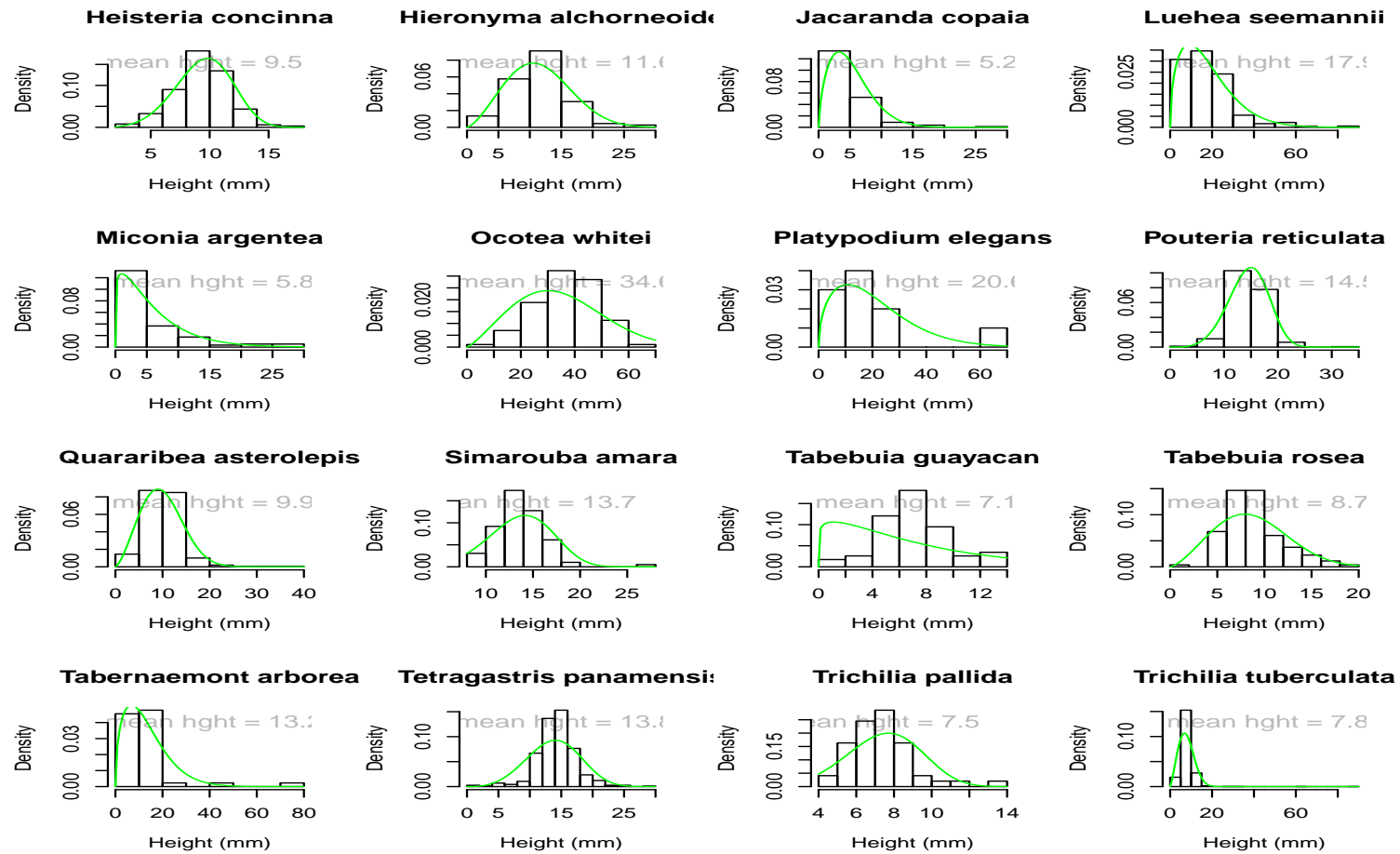


Figure S5. Distribution of initial heights of establishing seedlings, including fit models (green lines). Seedling heights of zero indicate very small seedlings that could not be measured without danger of inflicting damage. The above multi-panel graph is number 2 of 3 plots containing all species.

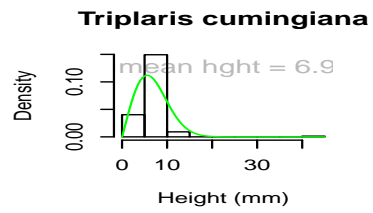


Figure S6. Distribution of initial heights of establishing seedlings, including fit models (green lines). Seedling heights of zero indicate very small seedlings that could not be measured without danger of inflicting damage. The above multi-panel graph is number 3 of 3 plots containing all species.

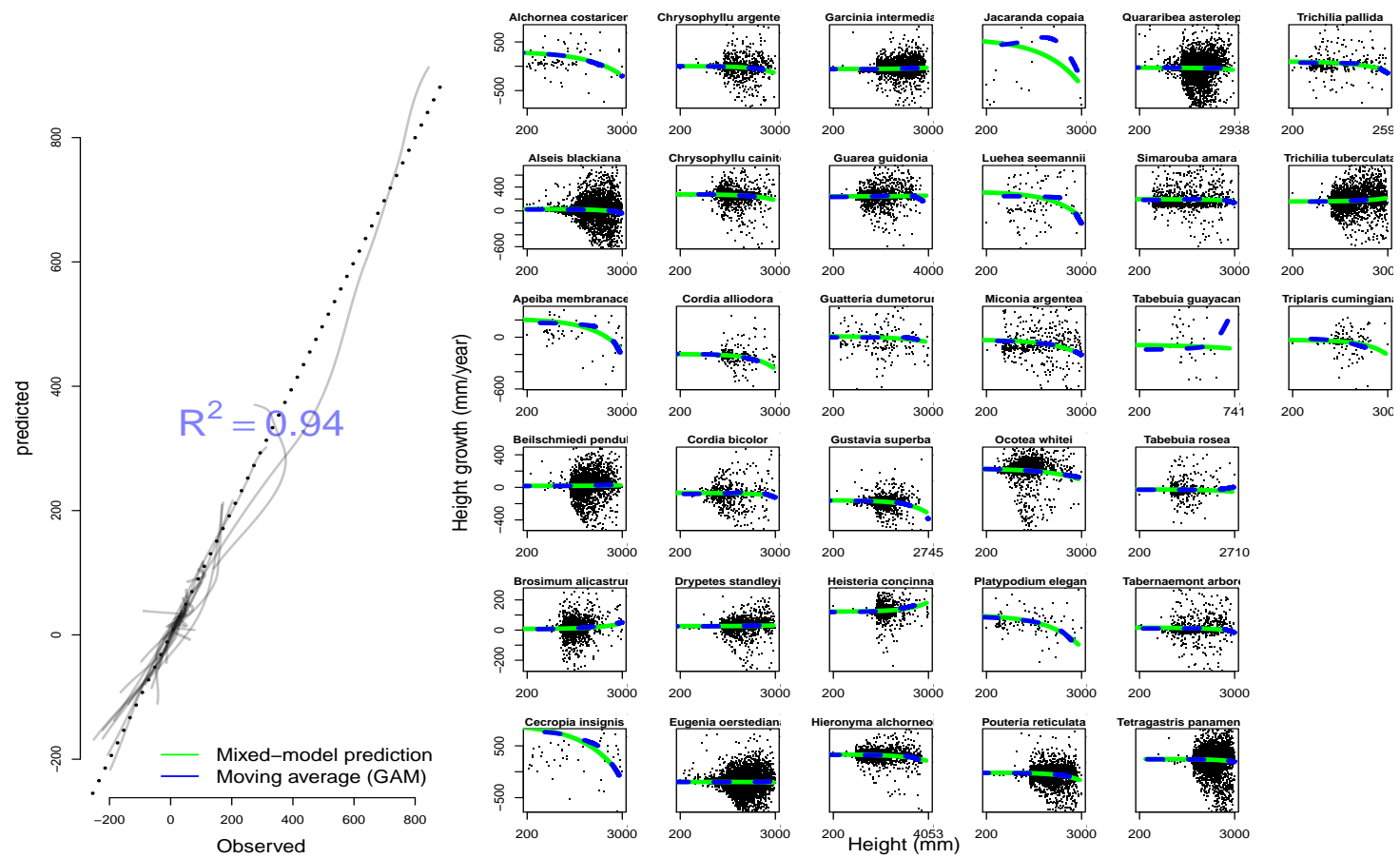


Figure S7. Yearly height growth of seedlings as a function of seedling height (mm). The large panel on the left shows the agreement between the model predictions (y-axis) from the mixed-model (see maintext) and an general additive model (GAM) fit for each species on the data (x-axis). The smaller panels on the right show model fits per species over size (height in mm). Here, the green line shows the predicted values by the fitted mixed effect model (corresponding to the y-axis on the large left panel), while the blue lines indicates a GAM predicted moving average (corresponding to the x-axis on the large left panel). The step-wise lines are indicative of the relative density of the data at a given size.

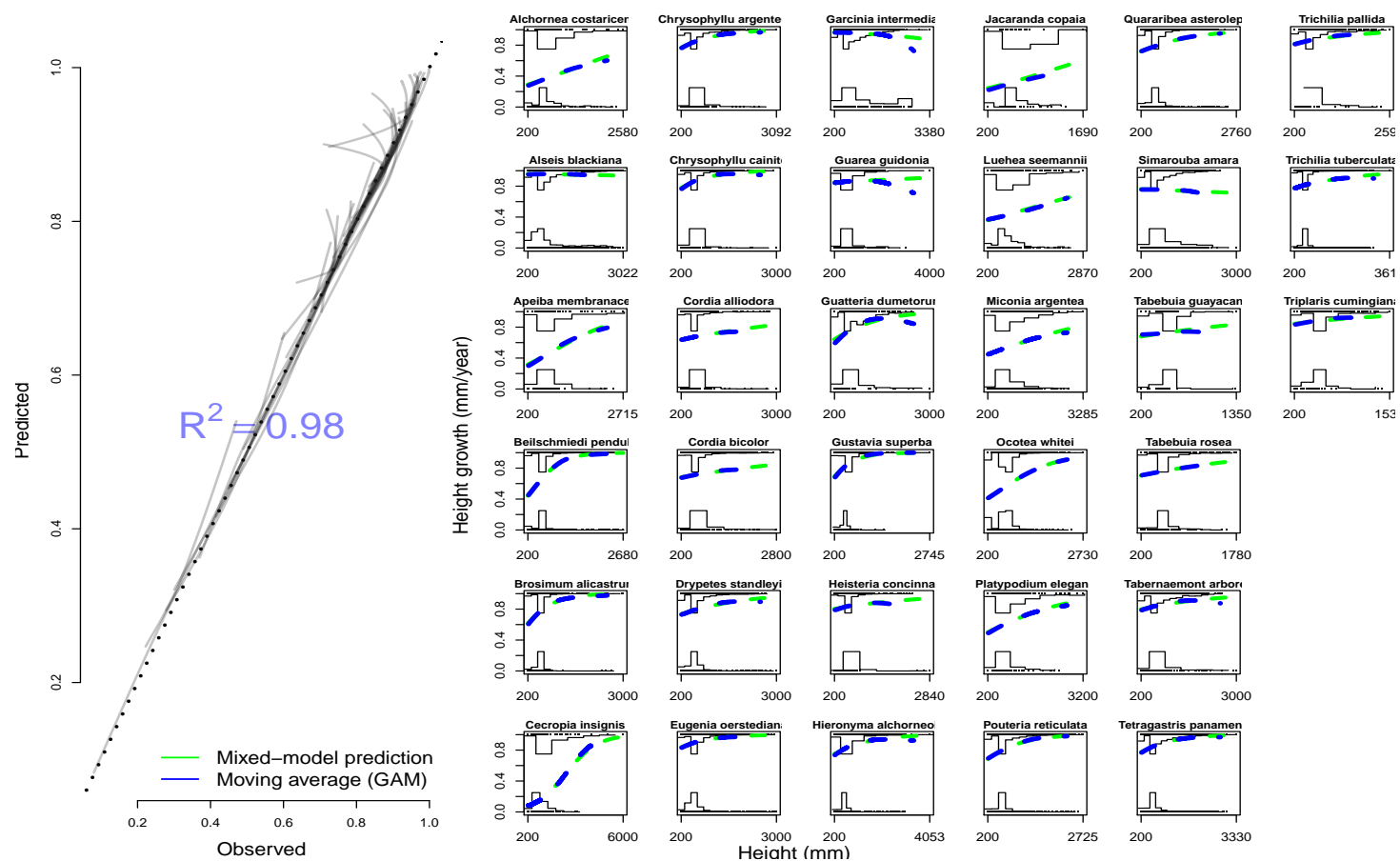


Figure S8. Yearly survival of seedlings as a function of seedling height (mm). The large panel on the left shows the agreement between the model predictions (y-axis) from the mixed-model (see maintext) and an general additive model (GAM) fit for each species on the data (x-axis). The smaller panels on the right show model fits per species over size (height in mm). Here, the green line shows the predicted values by the fitted mixed effect model (corresponding to the y-axis on the large left panel), while the blue lines indicates a GAM predicted moving average (corresponding to the x-axis on the large left panel). The step-wise lines are indicative of the relative density of the data at a given size. Individual survival and mortality states (1 and 0 values) are plotted as black dots that have been slightly jittered.

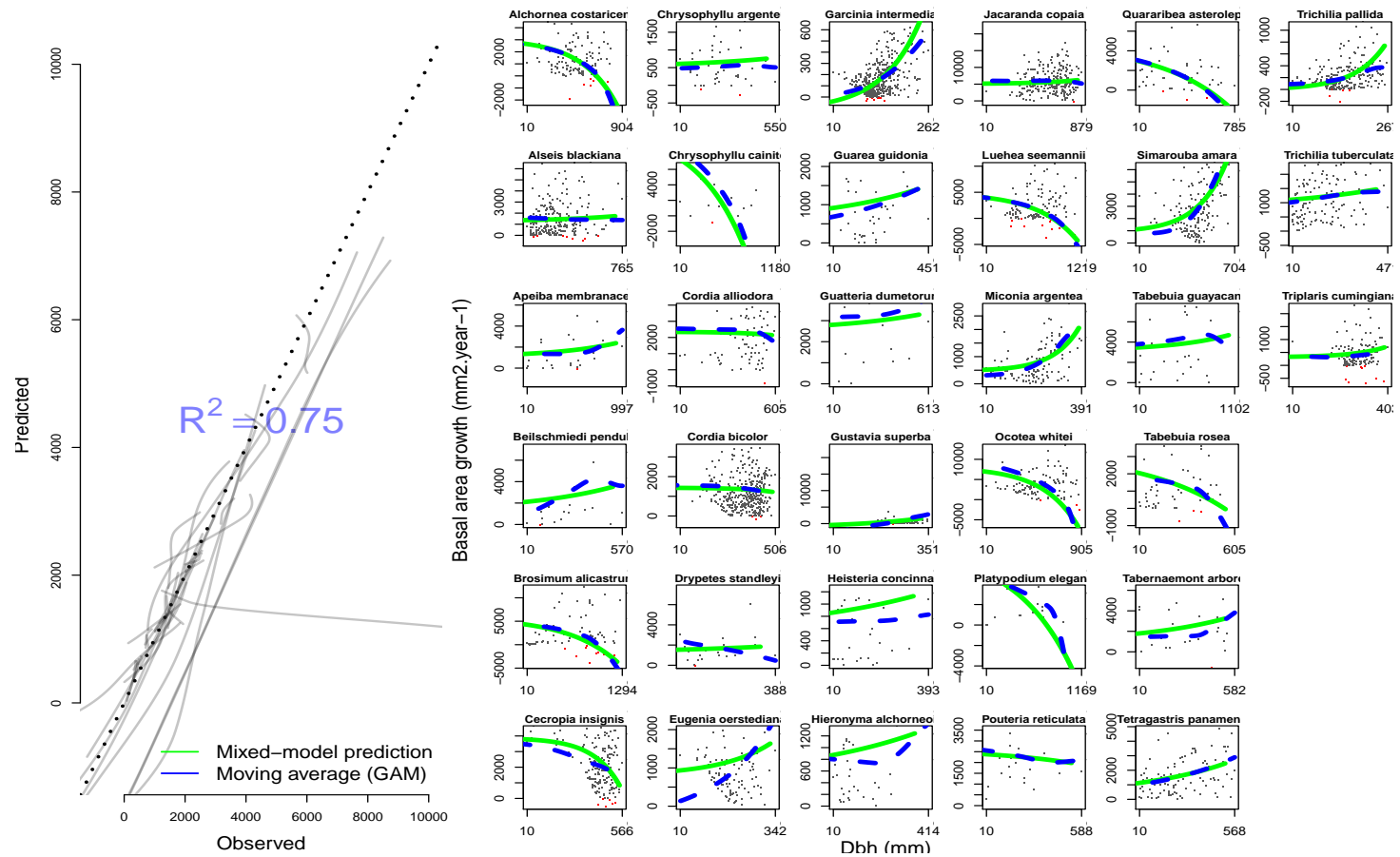


Figure S9. Yearly basal area growth of trees (mm^2), as a function of dbh (mm). The large panel on the left shows the agreement between the model predictions (y-axis) from the mixed-model (see maintext) and an general additive model (GAM) fit for each species on the data (x-axis). The smaller panels on the right show model fits per species over size (height in mm). Here, the green line shows the predicted values by the fitted mixed effect model (corresponding to the y-axis on the large left panel), while the blue lines indicates a GAM predicted moving average (corresponding to the x-axis on the large left panel).

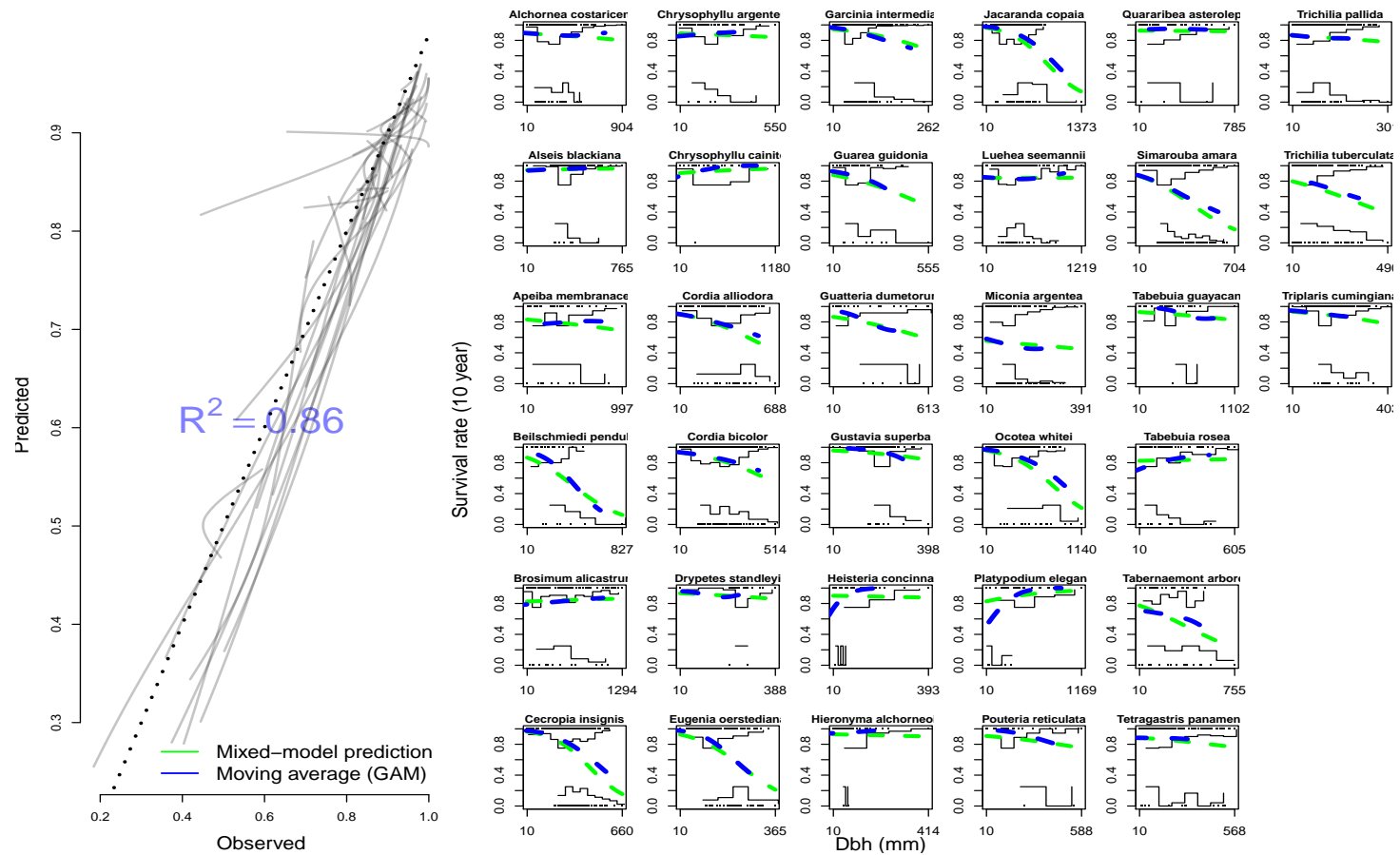


Figure S10. Decade survival of trees as a function of dbh (mm). The large panel on the left shows the agreement between the model predictions (y-axis) from the mixed-model (see maintext) and an general additive model (GAM) fit for each species on the data (x-axis). The smaller panels on the right show model fits per species over size (height in mm). Here, the green line shows the predicted values by the fitted mixed effect model (corresponding to the y-axis on the large left panel), while the blue lines indicates a GAM predicted moving average (corresponding to the x-axis on the large left panel). The stepwise lines show the relative density of mortality and survival events at different sizes classes. Individual survival and mortality states (1 and 0 values) are plotted as black dots that have been slightly jittered.

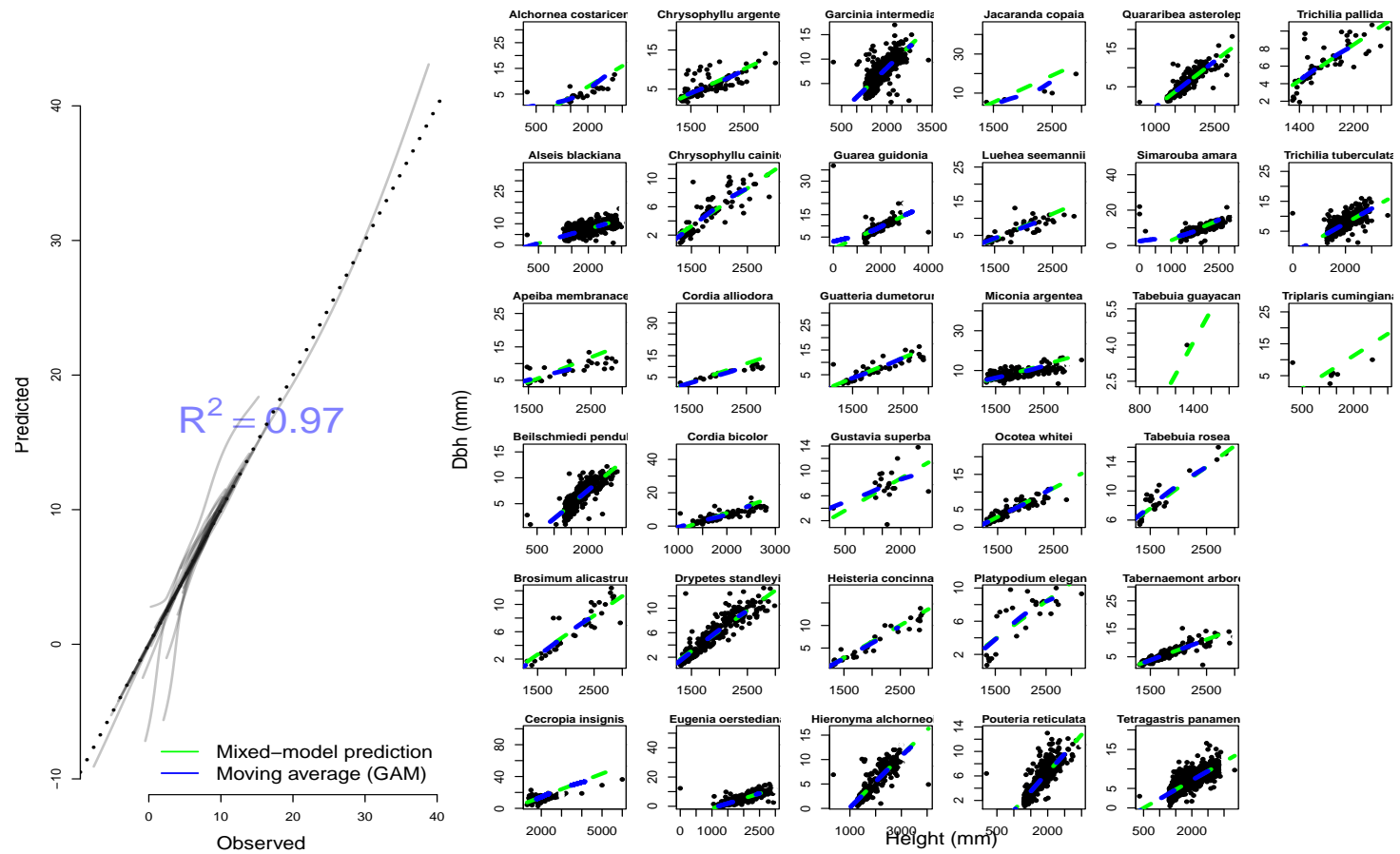


Figure S11. The diameter at breast height (dbh in mm) plotted against seedling height (mm), measured at the same time for each species. The large panel on the left shows the agreement between the model predictions (y-axis) from the mixed-model (see maintext) and an general additive model (GAM) fit for each species on the data (x-axis). The smaller panels on the right show model fits per species over size (height in mm). Here, the green line shows the predicted values by the fitted mixed effect model (corresponding to the y-axis on the large left panel), while the blue lines indicates a GAM predicted moving average (corresponding to the x-axis on the large left panel). Individual survival and mortality states (1 and 0 values) are plotted as black dots that have been slightly jittered.