

Alternative Balsam Fir Diameter Growth

Greg Johnson, David Marshall, Aaron Weiskittel

2026-01-28

Data

We extracted and processed Forest Inventory and Analysis (FIA) data from 10 states listed in the native range of Balsam Fir in the Silvics of North America.¹

After subsetting the data to censor observations with missing data, limiting the species to Balsam Fir (FIA species code 12), and remeasurement intervals ≥ 5 years we get the observations in Table 1.

Table 1: Balsam Fir Growth Observations by State

State	Observations
CT	1
MA	40
ME	83061
MI	24161
MN	40793
NH	11573
NY	7428
VT	4172
WI	22292
WV	6

¹Burns, Russell M., and Barbara H. Honkala, tech. coords. 1990. Silvics of North America: 1. Conifers; 2. Hardwoods. Agriculture Handbook 654. U.S. Department of Agriculture, Forest Service, Washington, DC. vol.2, 877 p.

Alternative Model Formulation

An alternative to the ORGANON diameter growth equation² which reduces parameter count while retaining key features of the original model is shown below. The key change is the term with a ratio of a transformation of diameter at breast height (**dbh**) squared to crown length. Since β_1 is expected to be negative, this tends to slow growth as more basal area accumulates in the tree while moderating that decline by the amount of productive crown capacity as measured by crown length. Basal area in larger trees (**bal**) serves as the inter-tree competition factor, and site index (**si**) as the inherent productivity scaling factor.

Site index is flawed for a number of reasons:

1. It is not consistently obtained for each plot due to missing Balsam Fir site trees,
2. It is estimated using a number of different and not necessarily compatible **si** equations, and
3. The available **si** equations do not all use the same base age.

In the data set **si** is derived from 27 different site index equations for 30 species. Balsam Fir site index comprises 33% of the observations. There are 1 base ages used. Preliminary graphical analysis revealed that base age was most correlated with residual bias. Thus in the following, we fit two equations: one where **SIBASE** and **SISP** are treated as a random effects in a mixed model framework, and a second leaving site index out.

$$\Delta dbh = e^{(\beta_0 + \beta_1 \log(\frac{(dbh+1)^2}{(cr*ht+1)^{\beta_4}}) + \beta_2 \frac{bal^{\beta_5}}{dbh+2.7} + \beta_3 \log(si_{s,b} + 4.5))} \quad (1)$$

and

$$\Delta dbh = e^{(\beta_0 + \beta_1 \log(\frac{(dbh+1)^2}{(cr*ht+1)^{\beta_4}}) + \beta_2 \frac{bal^{\beta_5}}{dbh+2.7})} \quad (2)$$

where:

- **dbh** = diameter at breast height (inches),
- **bal** = basal area per acre in larger trees ($feet^2/ac$),
- **cr** = crown ratio (fraction of total height),
- **ht** = total height (feet), and
- $si_{s,b}$ = site index (feet) for species **s** and base age **b**.
- $\beta_0 - \beta_5$ are parameters to be estimated.

²Hann, D.W., Marshall, D.D., and Hanus, M.L. 2006. Reanalysis of the SMC-ORGANON equations for diameter-growth rate, height-growth rate, and mortality rate of Douglas-fir. Forest Research Laboratory Research Contribution 49.

Nonlinear regression was used with an integrated fitting approach such that individual observations can have differing remeasurement intervals. The error to be minimized is ending dbh. Since this effectively minimizes diameter growth it can weight observations with longer remeasurement intervals more heavily. The effect of this needs to be evaluated, but putting more emphasis on longer periods may be beneficial.

The fit statistics for Equation 1 are:

Nonlinear mixed-effects model fit by maximum likelihood

Model: endDIA ~ est_dg(B0, B1, B2, B3, B4, B5, startDIA, startBAL, endBAL, startCR, endCR)

Data: tree_subset %>% mutate(SIINT = interaction(as.factor(tree_subset\$SIBASE), as.factor(tree_subset\$SIBASE)))

	AIC	BIC	logLik
	105104	105185.4	-52543.99

Random effects:

Formula: B3 ~ 1 | SIINT

	B3	Residual
StdDev:	0.01097501	0.3174408

StdDev: 0.01097501 0.3174408

Fixed effects: B0 + B1 + B2 + B3 + B4 + B5 ~ 1

	Value	Std.Error	DF	t-value	p-value
B0	-3.494068	0.029842193	193520	-117.08482	0
B1	-0.414431	0.003233631	193520	-128.16264	0
B2	-0.068861	0.005162066	193520	-13.33987	0
B3	0.088204	0.010242857	193520	8.61125	0
B4	2.356021	0.014610190	193520	161.25874	0
B5	0.609137	0.013398577	193520	45.46285	0

Correlation:

	B0	B1	B2	B3	B4
B1	-0.243				
B2	-0.402	0.528			
B3	-0.523	-0.008	-0.038		
B4	-0.405	0.774	0.655	-0.089	
B5	-0.377	0.499	0.994	-0.031	0.599

Standardized Within-Group Residuals:

	Min	Q1	Med	Q3	Max
	-31.9347814	-0.5754987	-0.1648738	0.4248148	24.6460943

Number of Observations: 193527

Number of Groups: 2

\$SIINT

B3
50.FALSE -0.01096438
50.TRUE 0.01096438

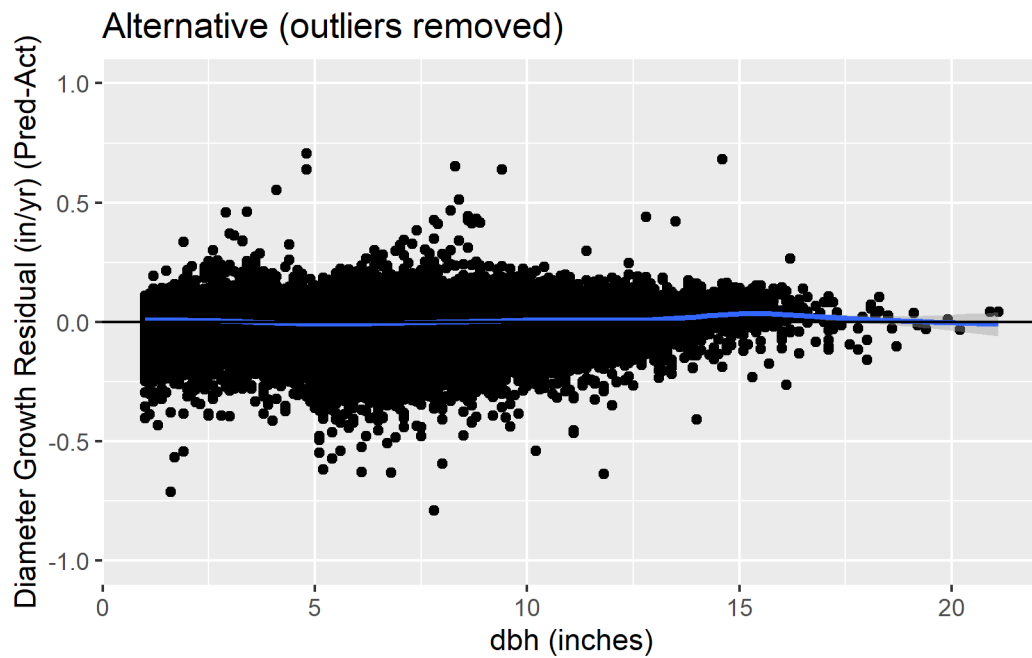
Residual Standard Error: 0.317440841300731 on 193520 degrees of freedom, AIC: 105104

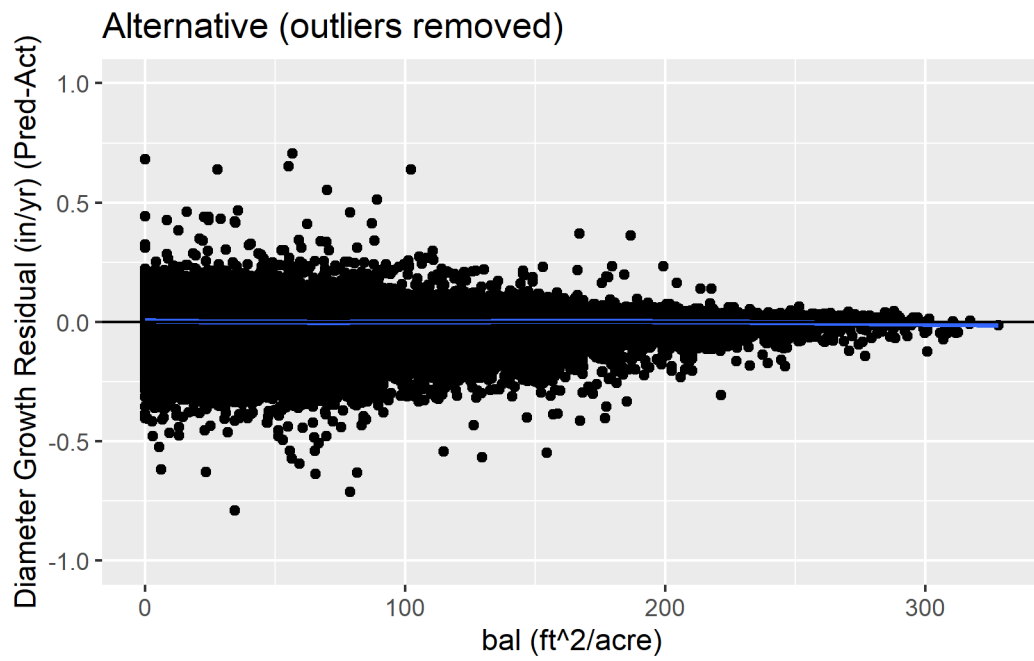
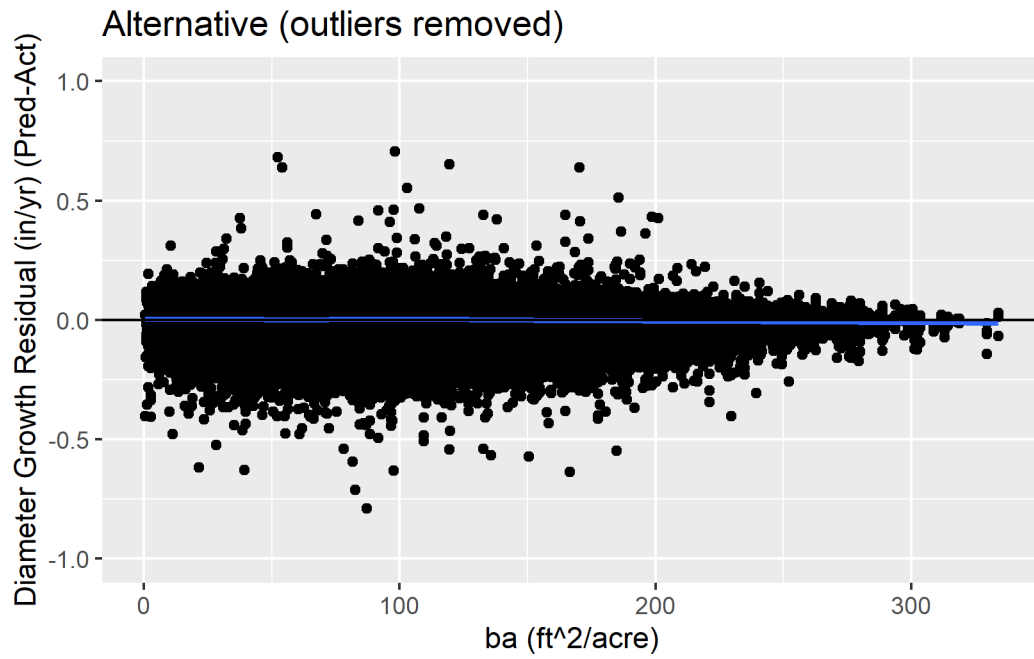
and for Equation 2:

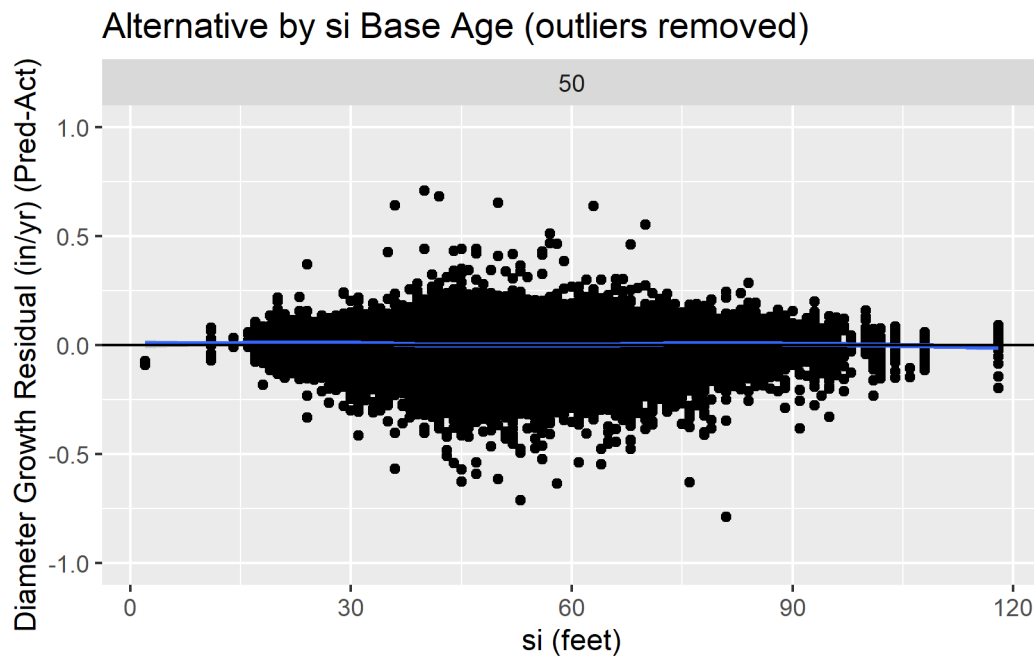
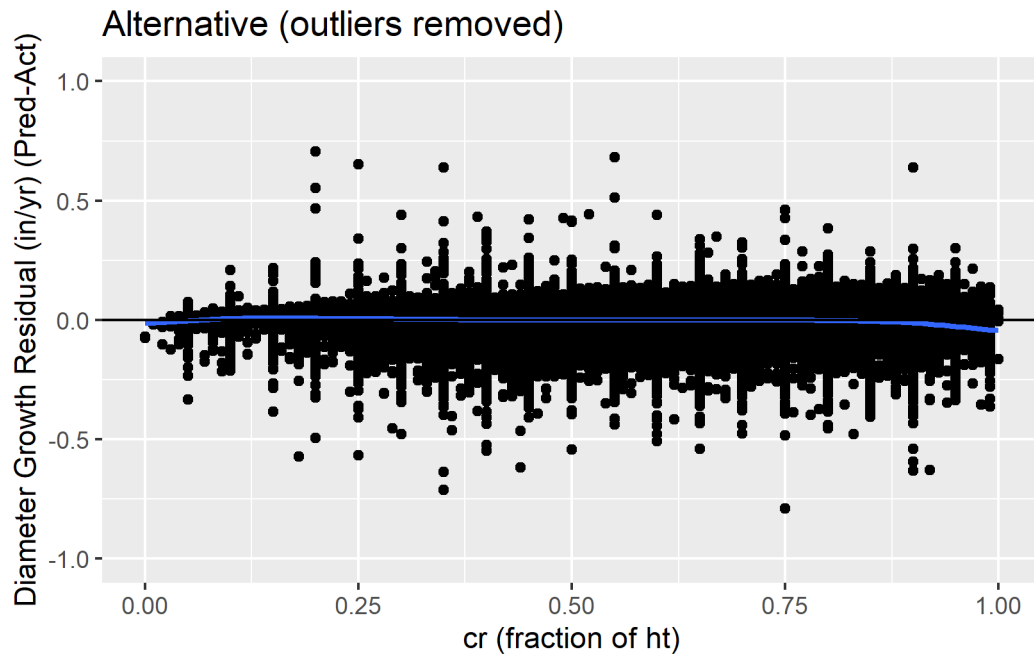
	Coef.	Std. error	t-stat.	p
B0	-3.1635990	0.0176928	-178.80727	0
B1	-0.4015284	0.0031861	-126.02518	0
B2	-0.0650773	0.0048980	-13.28642	0
B4	2.3993930	0.0151128	158.76598	0
B5	0.6206128	0.0134995	45.97285	0

Residual Standard Error: 0.318189813937393 on 193522 degrees of freedom, AIC: 105994.5

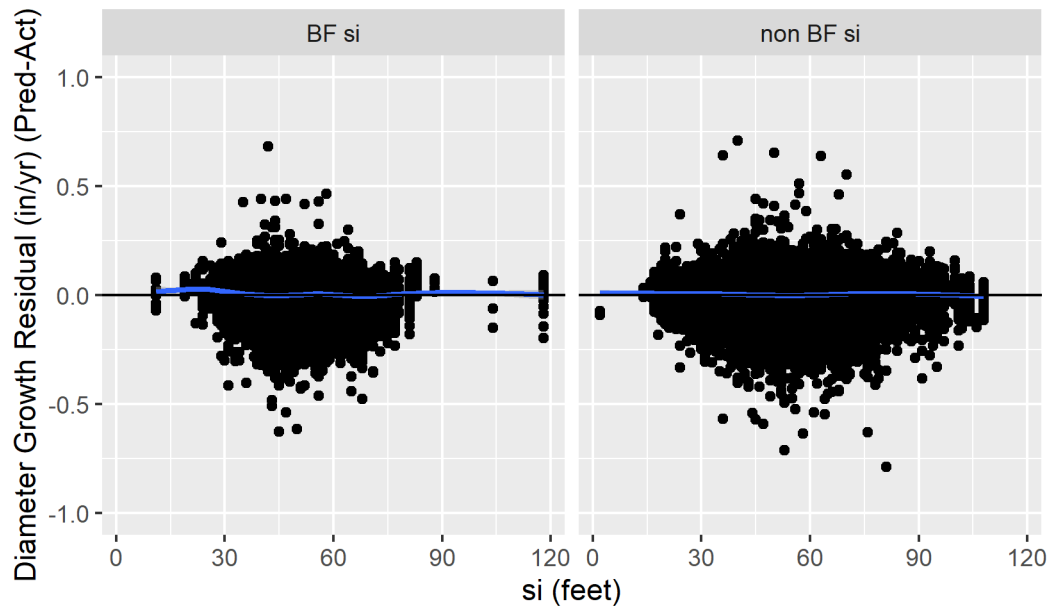
Residual Analysis for Equation 1



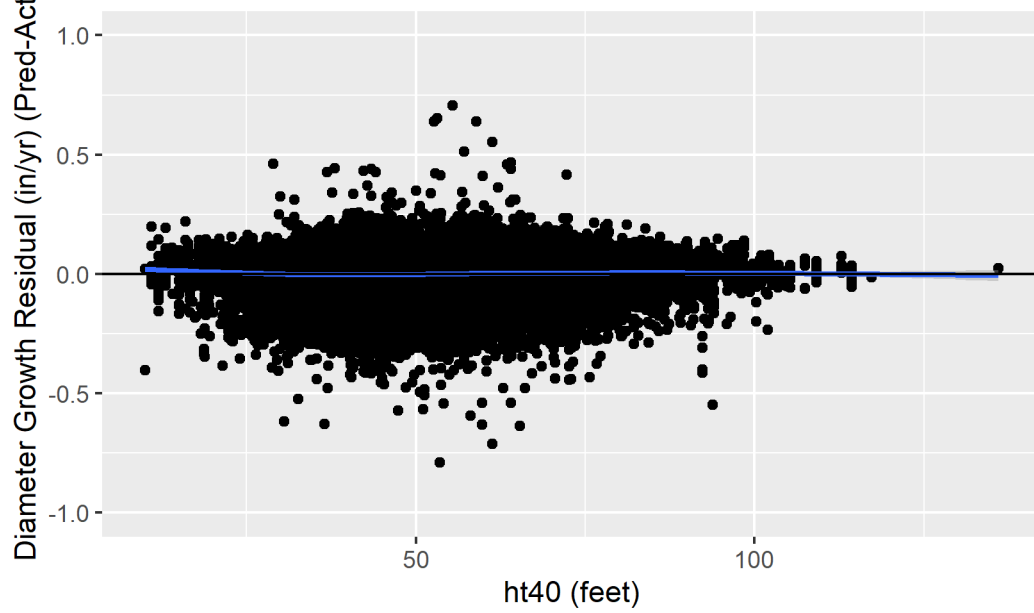


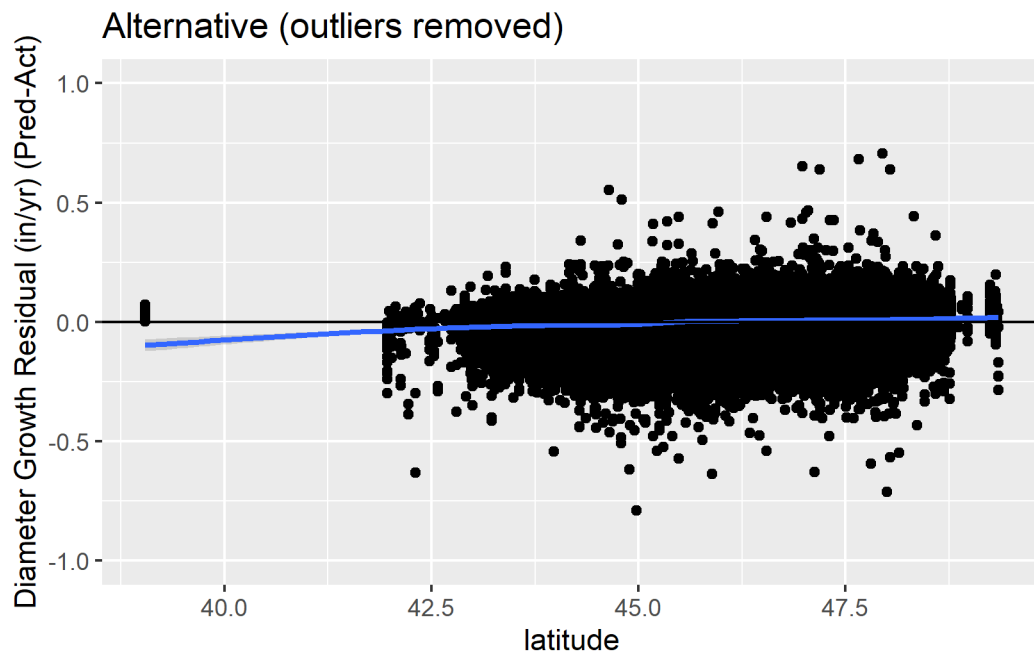
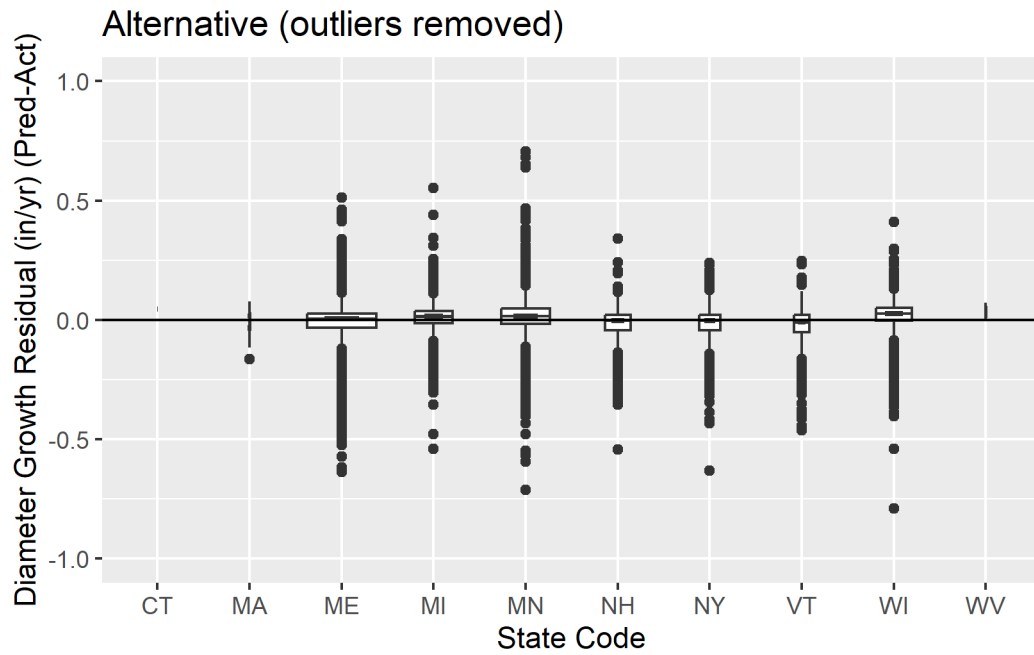


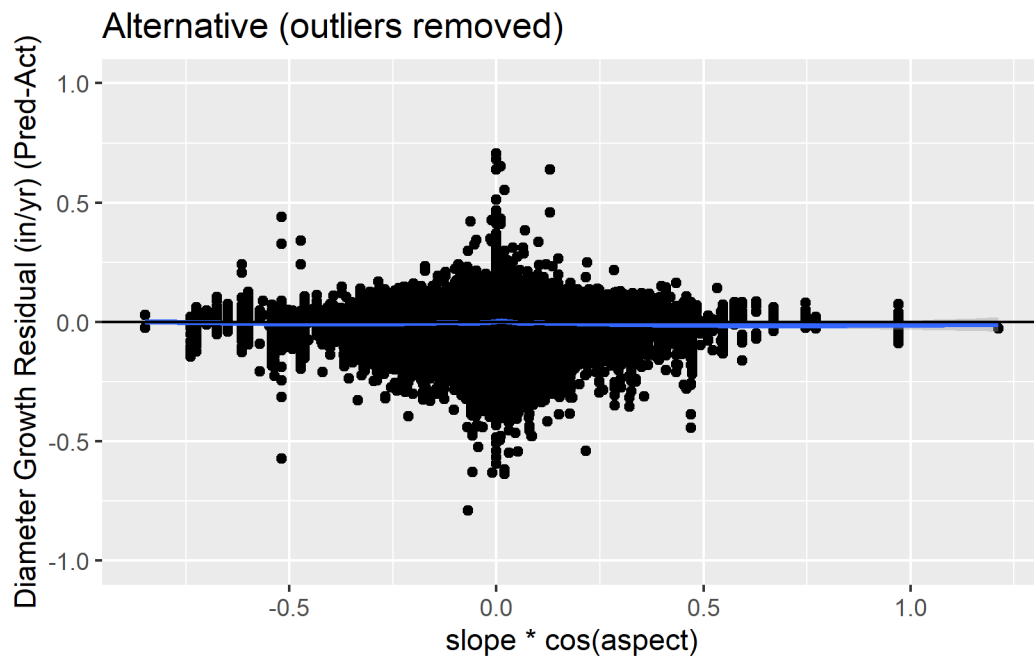
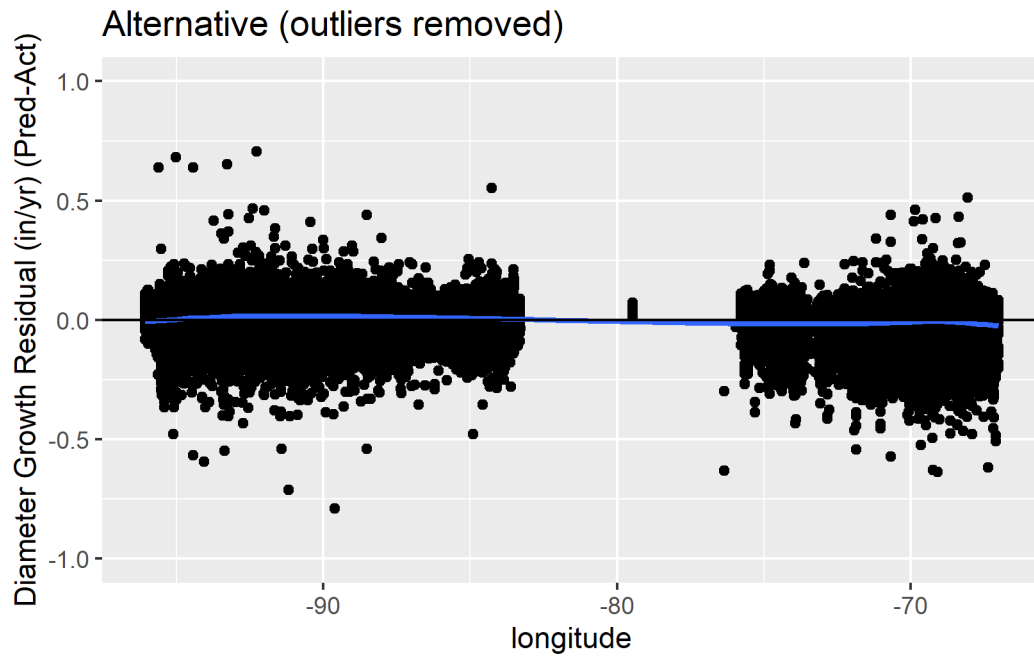
Alternative by si Species (outliers removed)

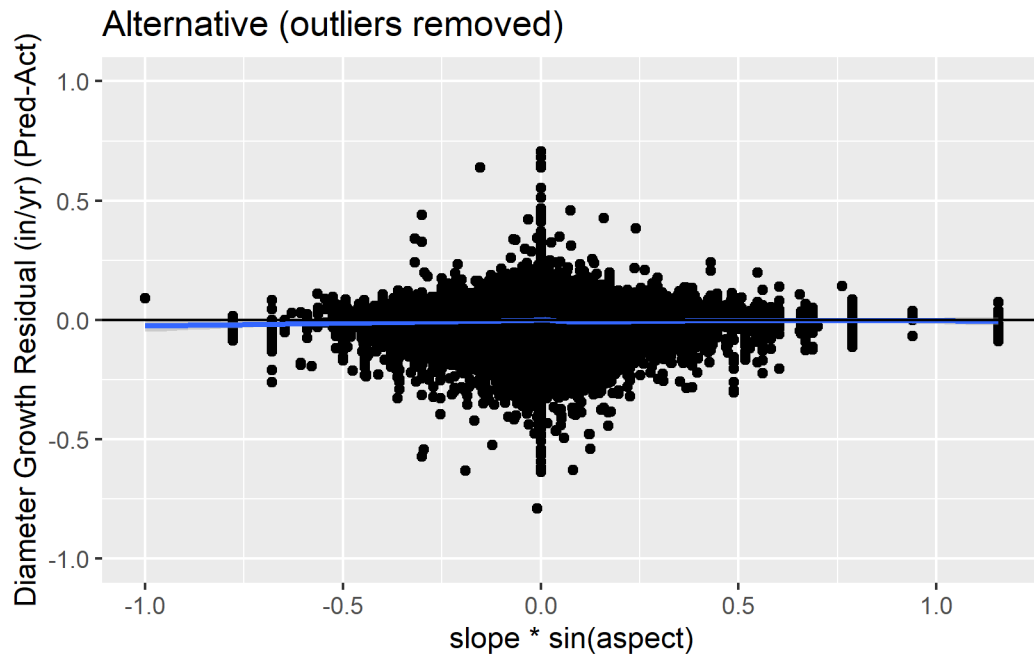


Alternative (outliers removed)

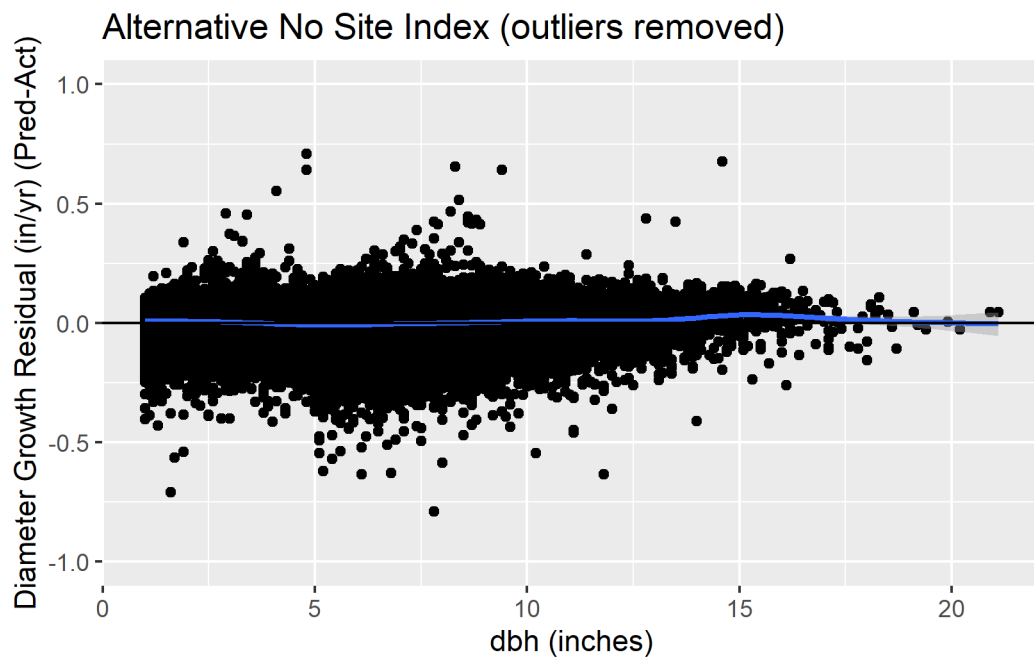


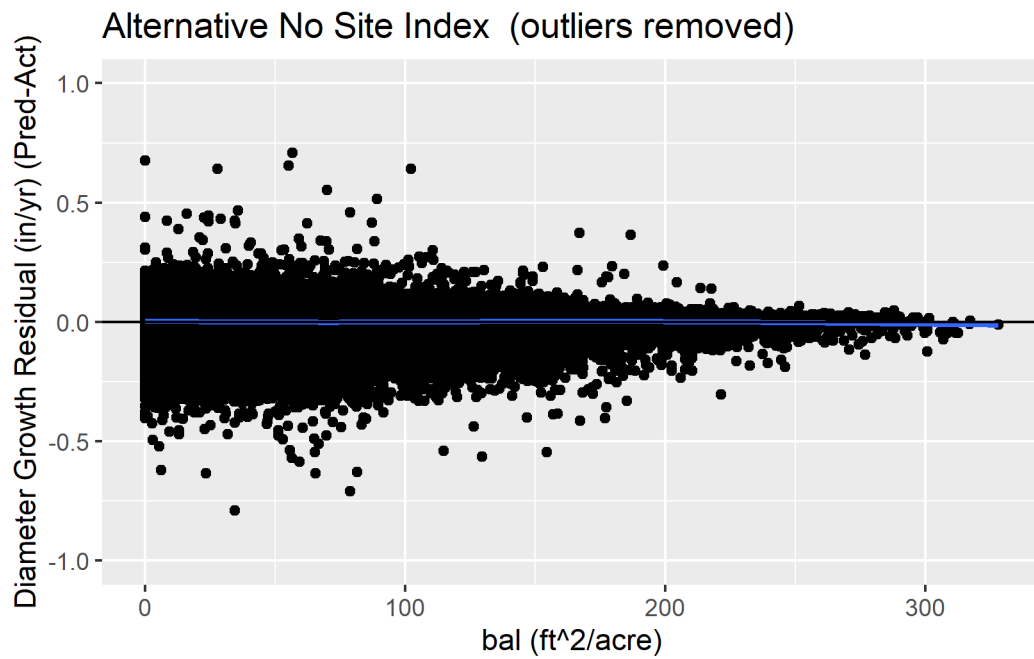
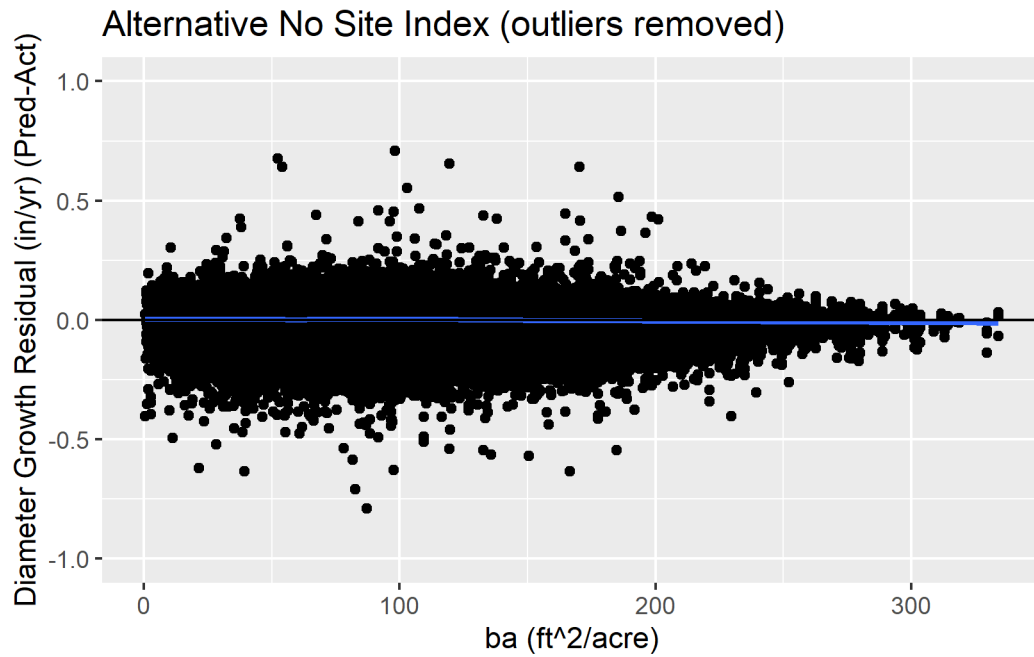


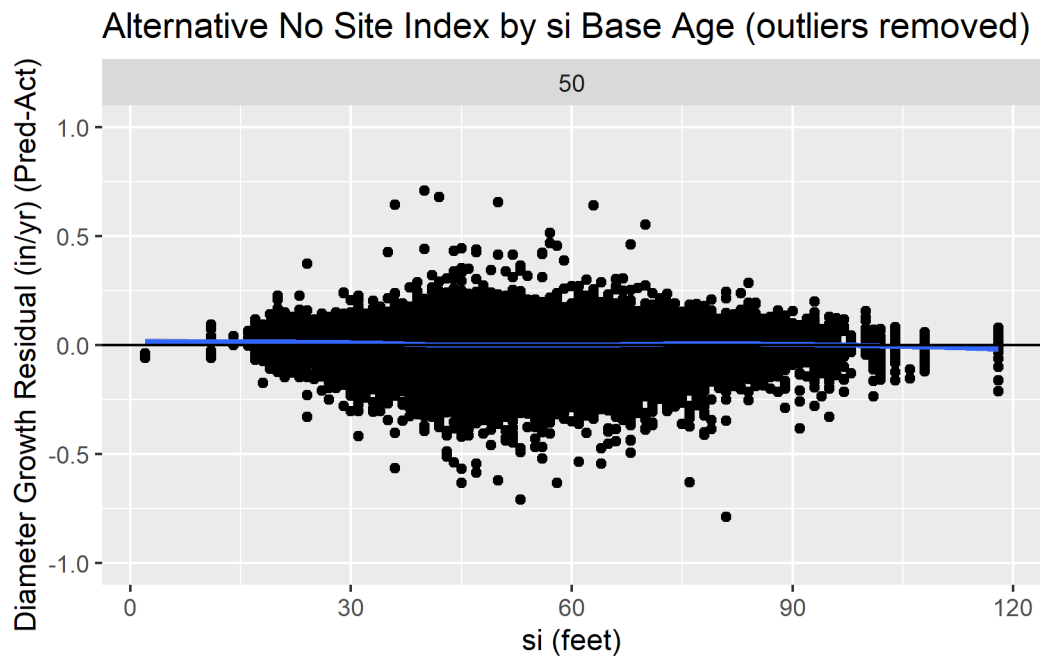
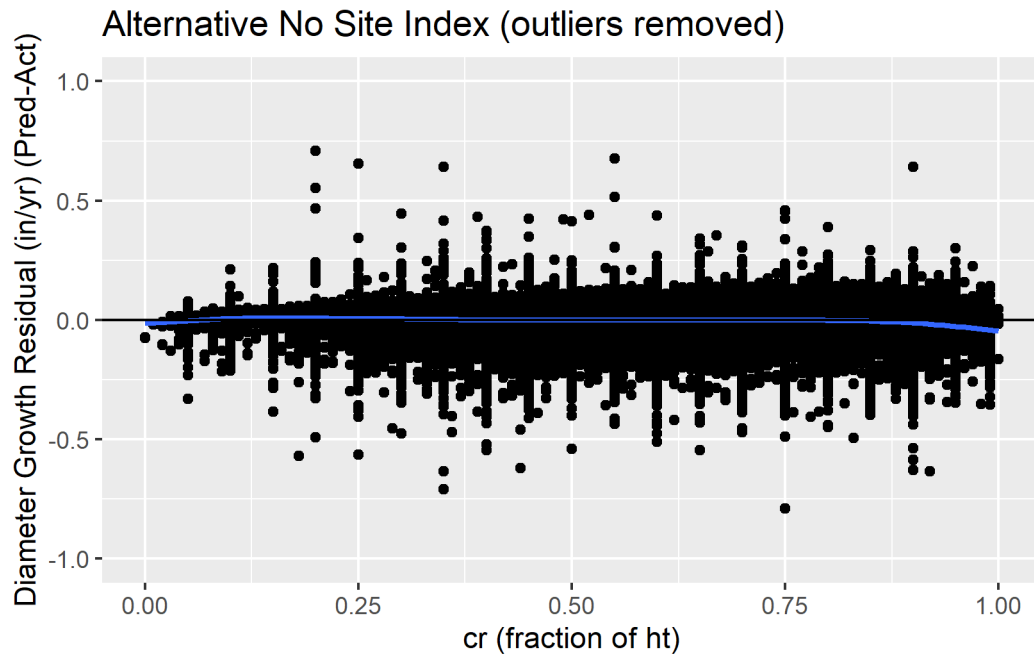




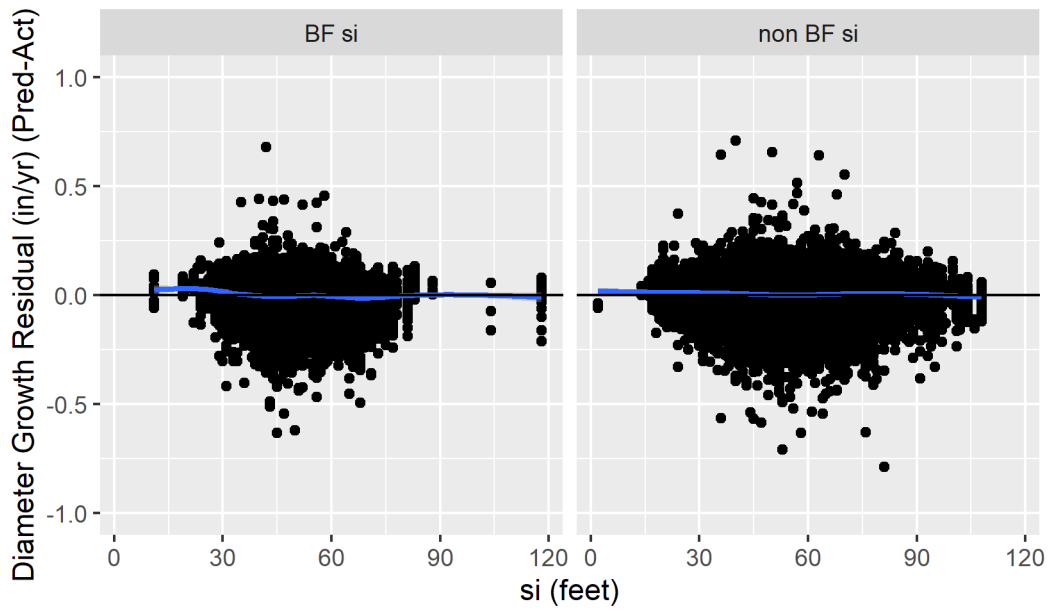
Residual Analysis for Equation 2



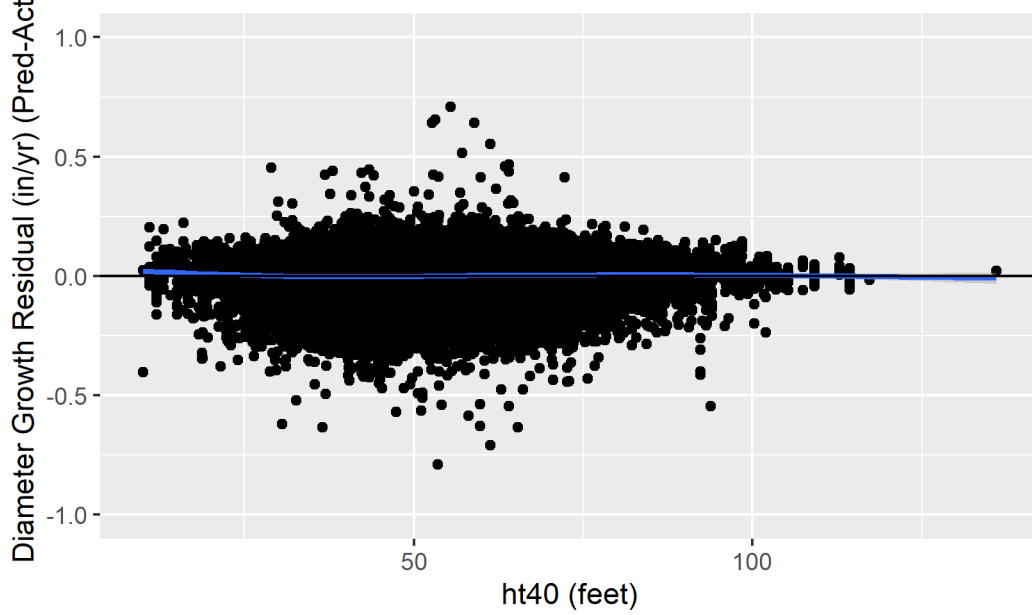


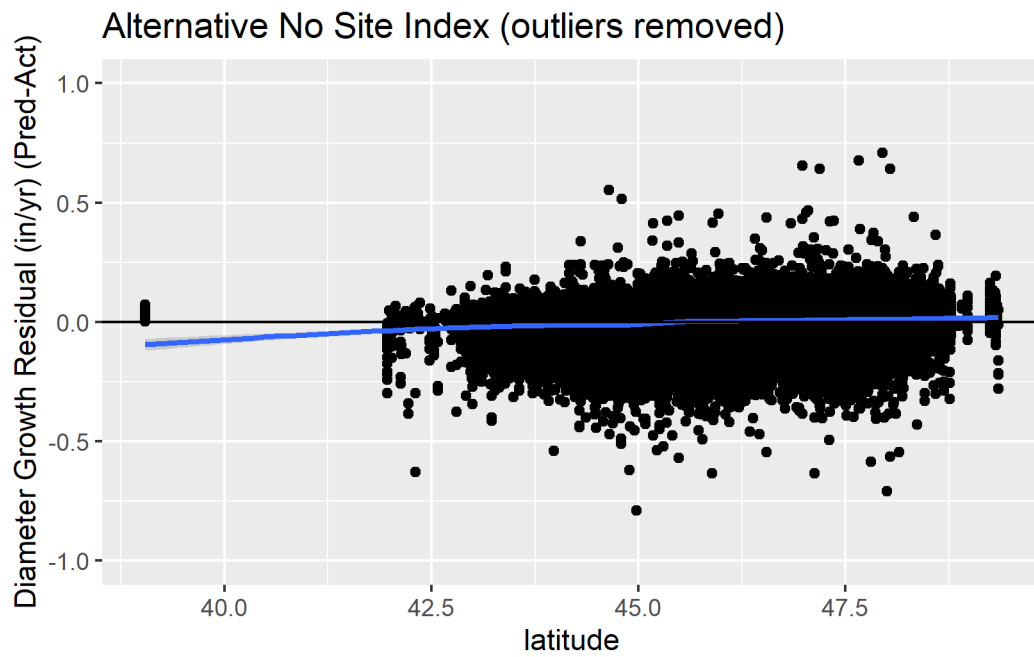
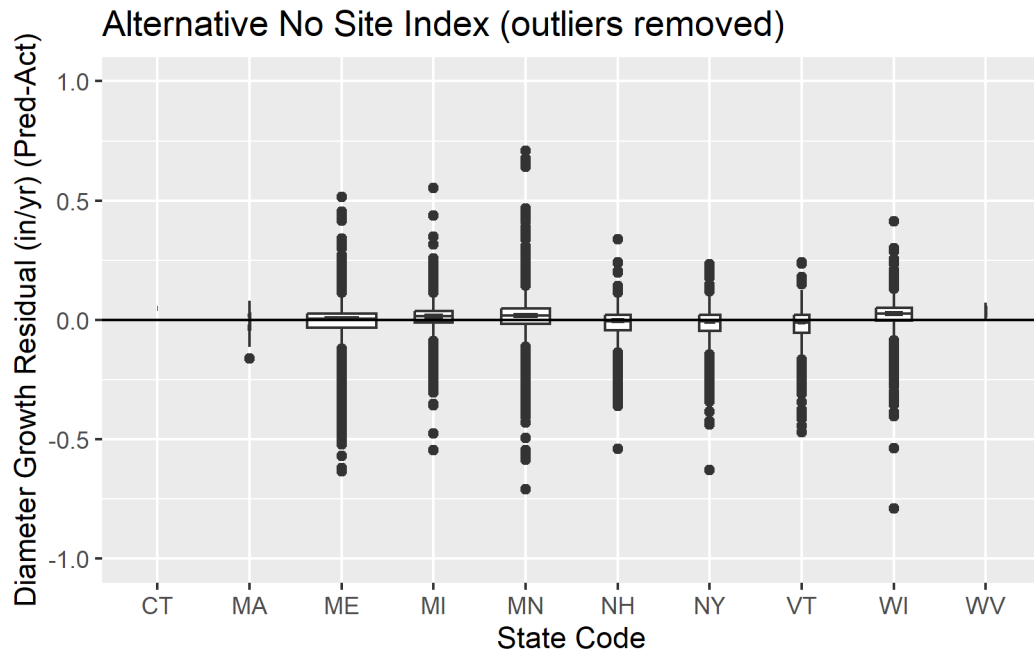


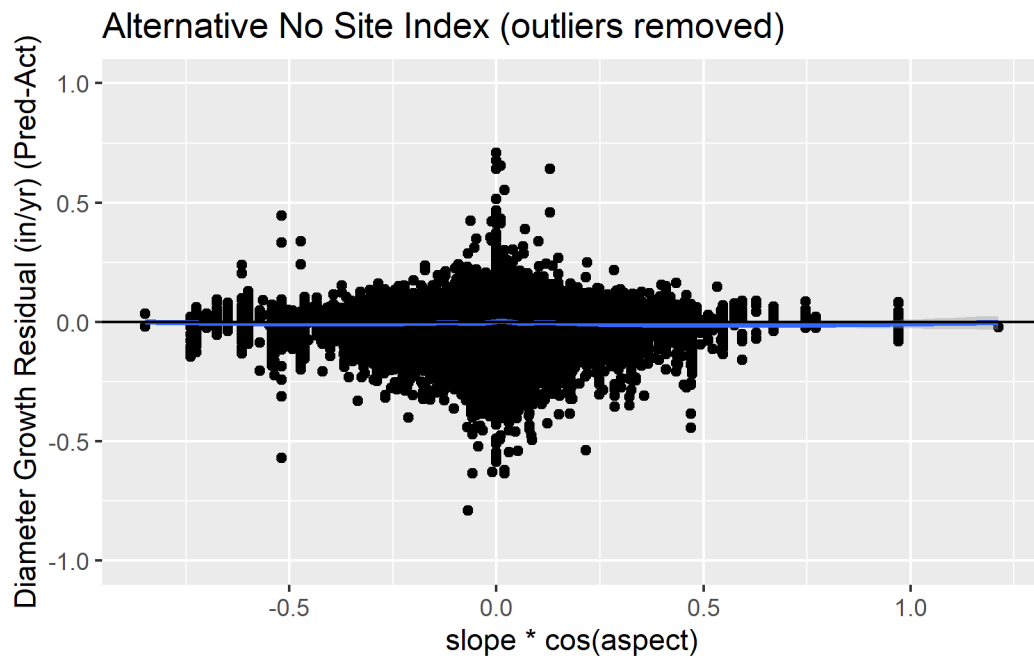
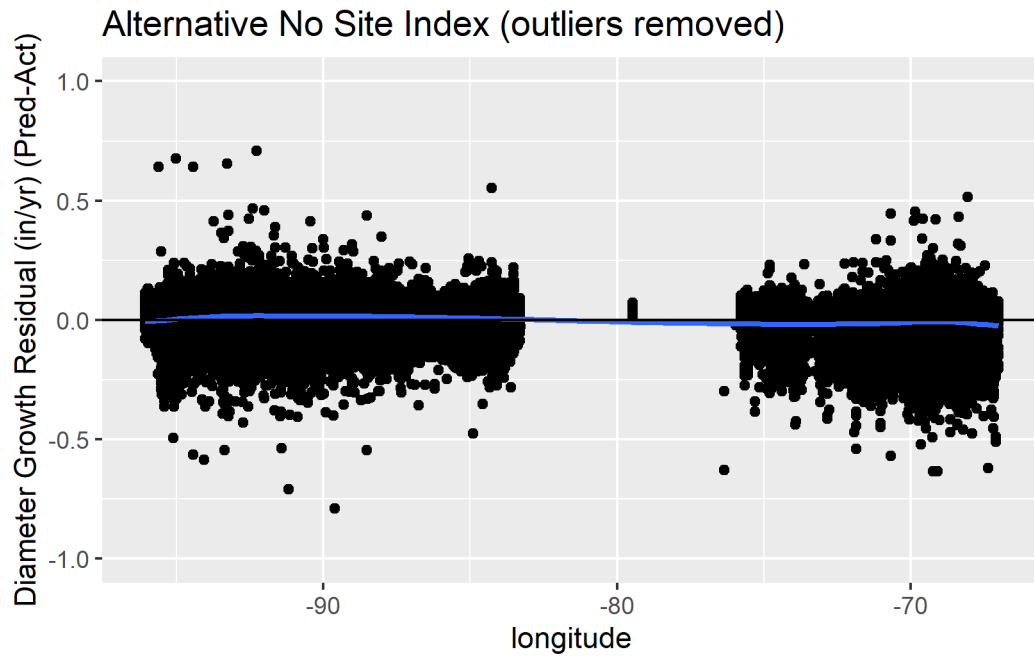
Alternative No Site Index by si Species (outliers removed)

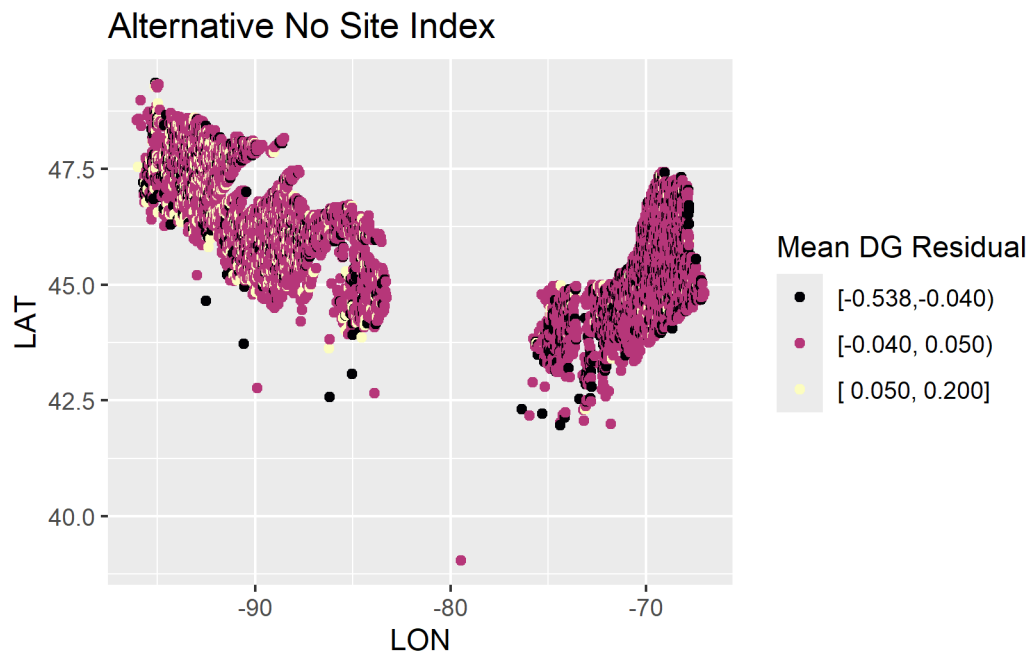
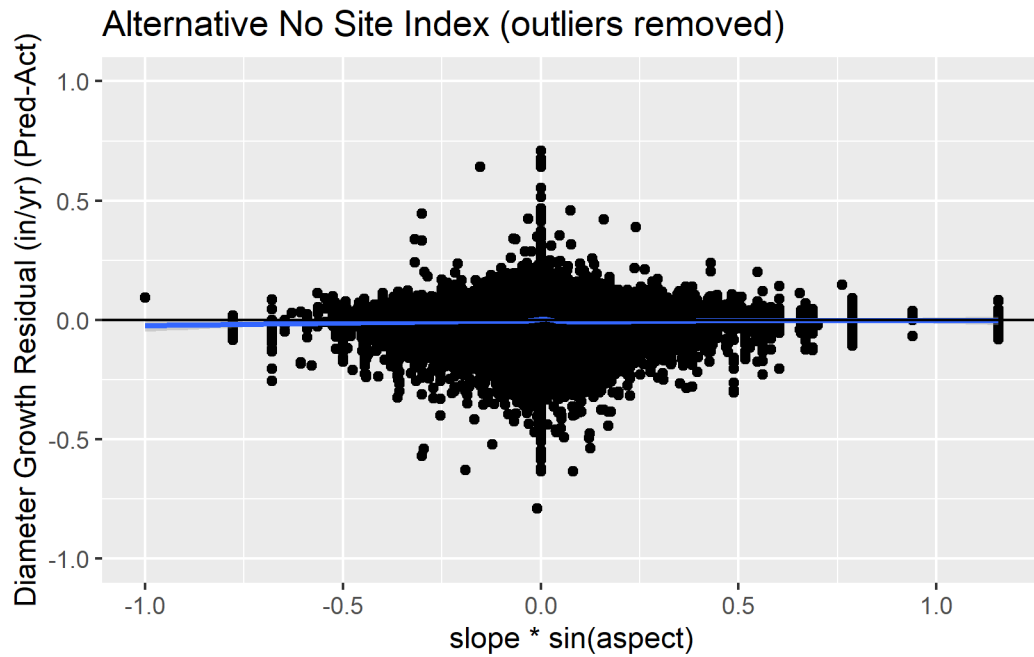


Alternative No Site Index (outliers removed)









Discussion

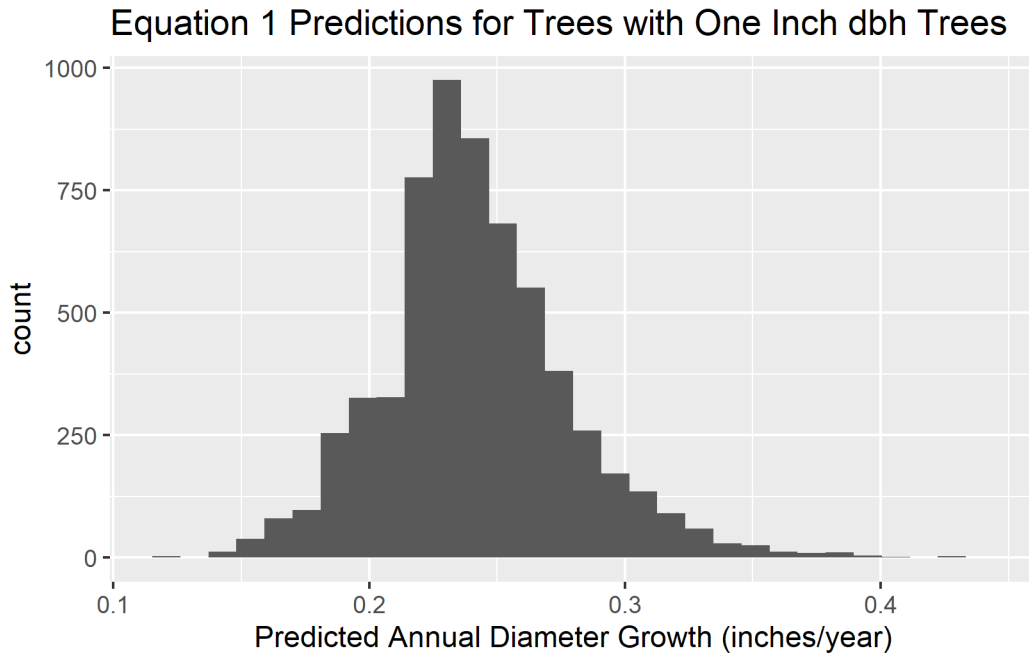
Removing `si` does not degrade the fit significantly. Only MN appears to be over-predicted.

Table 3: Independent Variables for One Inch dbh Trees

Variable	N	Mean	Std. Dev.	Min	Pctl. 25	Pctl. 75	Max
ba	6185	111	49	0.41	76	145	313
bal	6185	110	49	0	74	144	313
ht	6185	9.9	0.91	5	10	10	25
cr	6185	0.48	0.22	0.03	0.3	0.65	1
si	6185	51	13	17	44	59	108

Equation 1 residual graphs show that there is some indication of residual trends spatially (especially in the western states).

Equation Behavior for Very Small Trees



Equation 2 Predictions for Trees with One Inch dbh Trees

