

Alternative Incense Cedar Diameter Growth

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Data

We extracted and processed Forest Inventory and Analysis (FIA) data from 3 states listed in the native range of Incense Cedar in the Silvics of North America.¹

After subsetting the data to censor observations with missing data, limiting the species to Incense Cedar (FIA species code 81), and remeasurement intervals ≥ 5 years we get the observations in Table 1.

Table 1: Incense Cedar Growth Observations by State

State	Observations
CA	7314
NV	3
OR	2615

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

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.

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.

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 Incense Cedar 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 17 different site index equations for 17 species. Incense Cedar site index comprises 1% of the observations. There are 2 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 (*feet²/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.

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$SIBASE), as.factor(tree_subset$SISP)))
AIC      BIC      logLik
```

21633.61 21691.24 -10808.8

Random effects:

Formula: B3 ~ 1 | SIINT

B3 Residual

StdDev: 0.0632119 0.7176832

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

	Value	Std.Error	DF	t-value	p-value
B0	-4.658636	0.13899846	9923	-33.51574	0
B1	-0.694860	0.01438152	9923	-48.31613	0
B2	-0.149961	0.03570928	9923	-4.19950	0
B3	0.601195	0.04117971	9923	14.59930	0
B4	1.783763	0.02826586	9923	63.10665	0
B5	0.521112	0.03820534	9923	13.63976	0

Correlation:

	B0	B1	B2	B3	B4
B1	-0.314				
B2	-0.527	0.533			
B3	-0.475	0.032	0.014		
B4	-0.522	0.415	0.626	-0.095	
B5	-0.498	0.494	0.991	0.027	0.550

Standardized Within-Group Residuals:

	Min	Q1	Med	Q3	Max
	-9.51836151	-0.50950413	-0.09309793	0.40641442	29.30518247

Number of Observations: 9932

Number of Groups: 4

\$SIINT

	B3
50.FALSE	0.07974169
100.FALSE	-0.02274200
50.TRUE	0.03159499
100.TRUE	-0.08859469

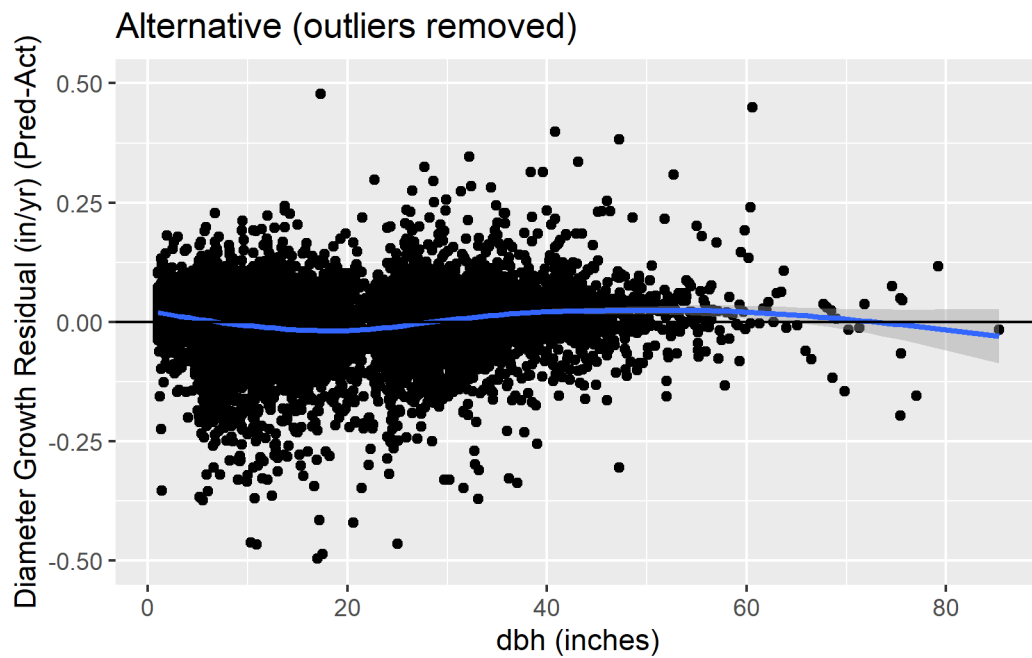
Residual Standard Error: 0.717683216123872 on 9923 degrees of freedom, AIC: 21633.6

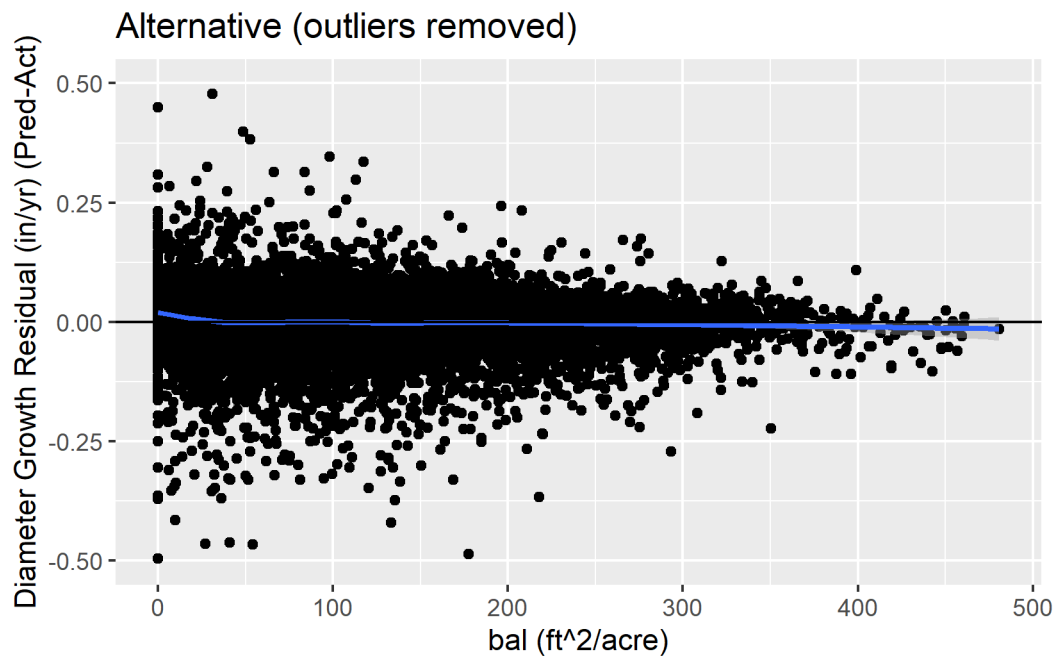
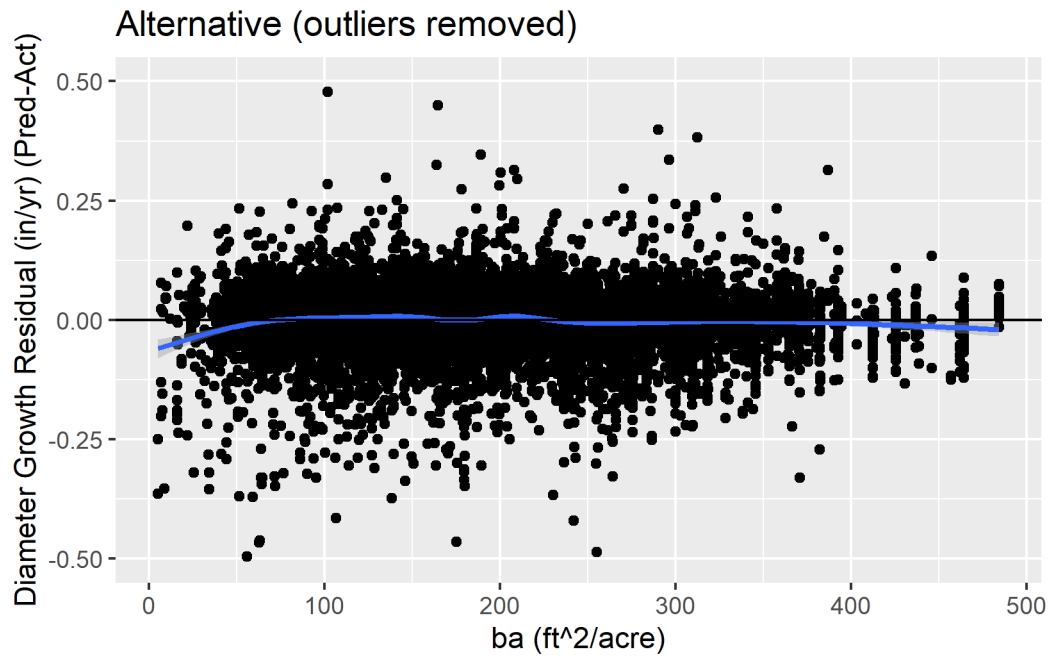
and for Equation 2:

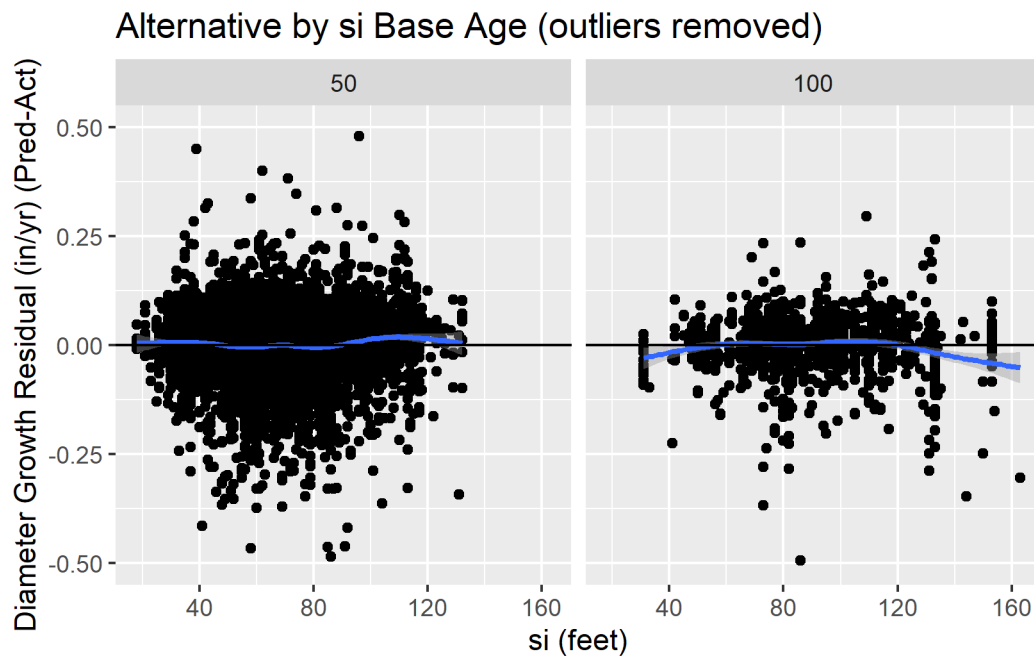
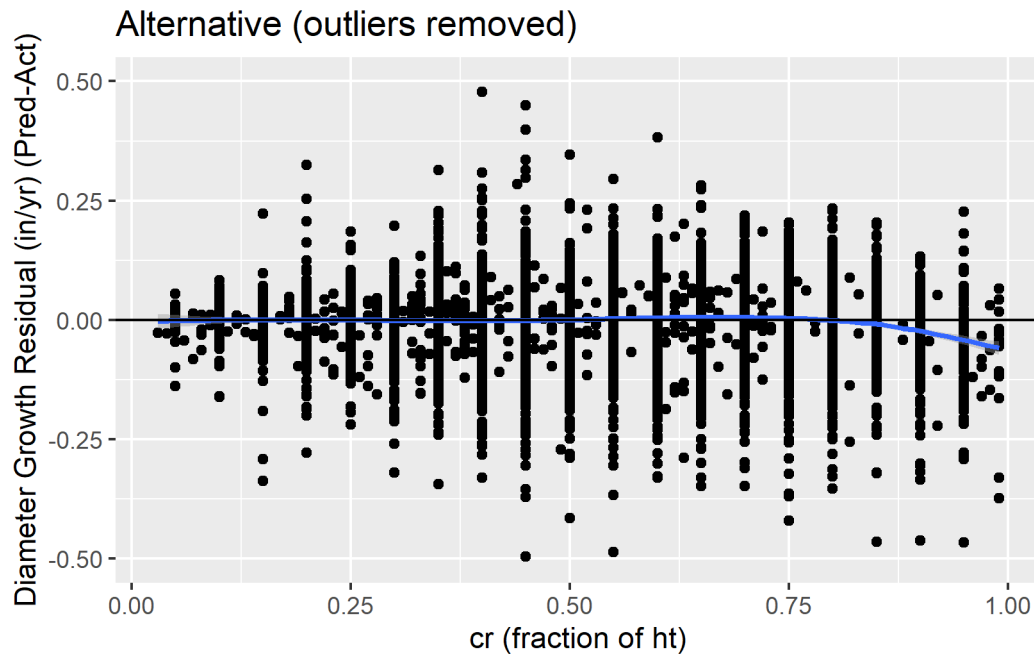
	Coef.	Std. error	t-stat.	p
B0	-2.5198137	0.0992009	-25.401110	0.0000000
B1	-0.6838018	0.0150817	-45.339705	0.0000000
B2	-0.1342665	0.0440084	-3.050933	0.0022873
B4	1.9950859	0.0319159	62.510633	0.0000000
B5	0.4830347	0.0517938	9.326119	0.0000000

Residual Standard Error: 0.744315396278852 on 9927 degrees of freedom, AIC: 22327.1

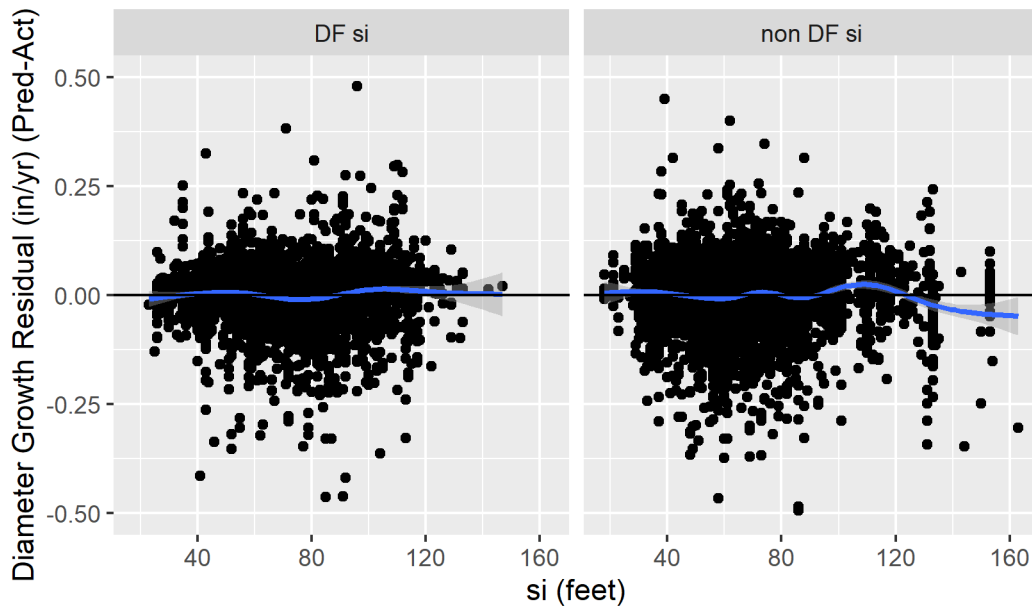
Residual Analysis for Equation 1



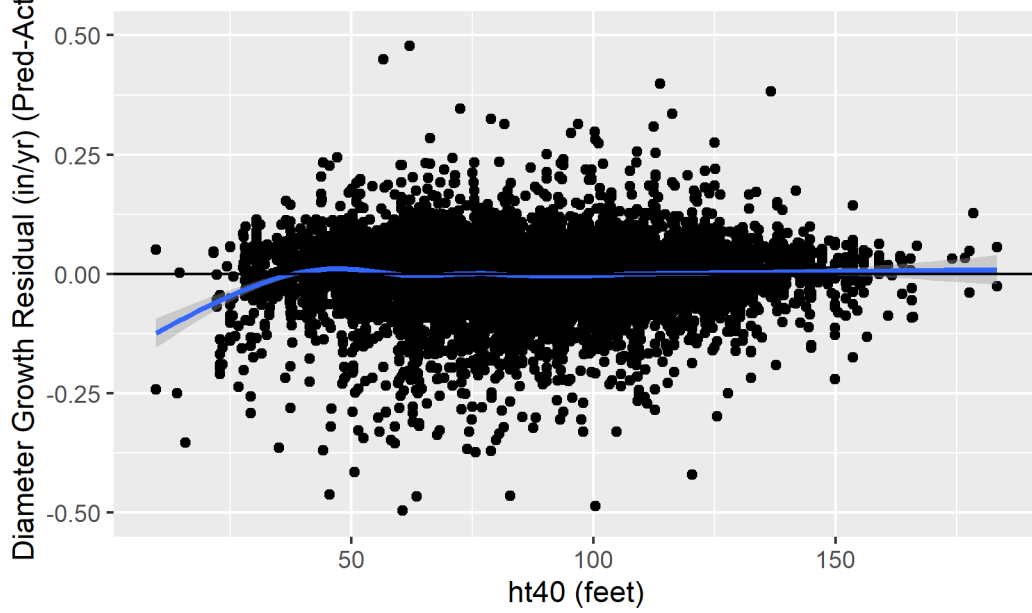


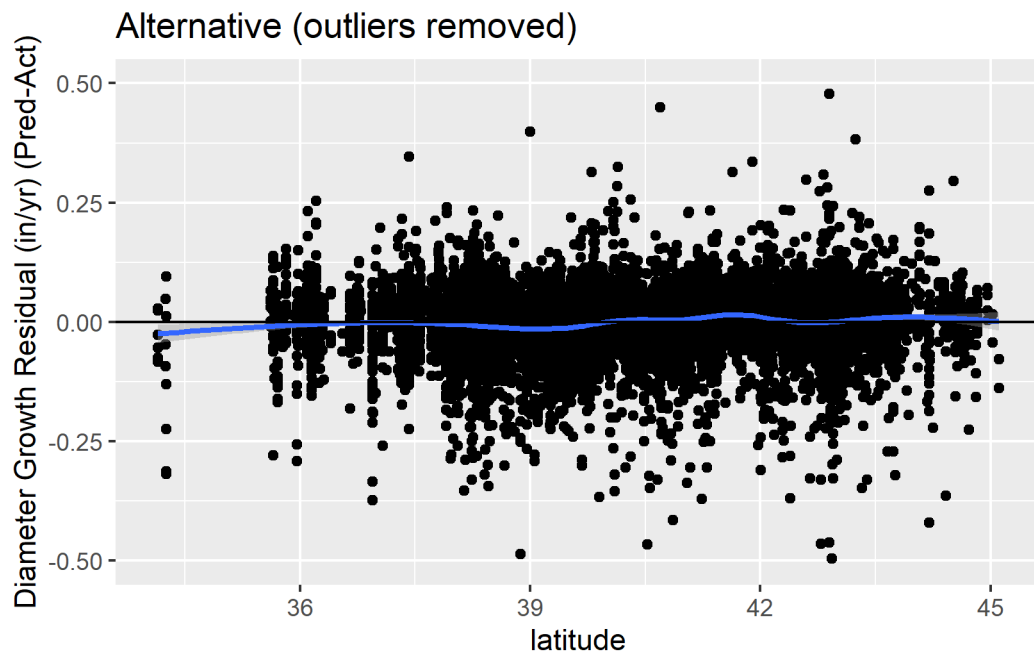
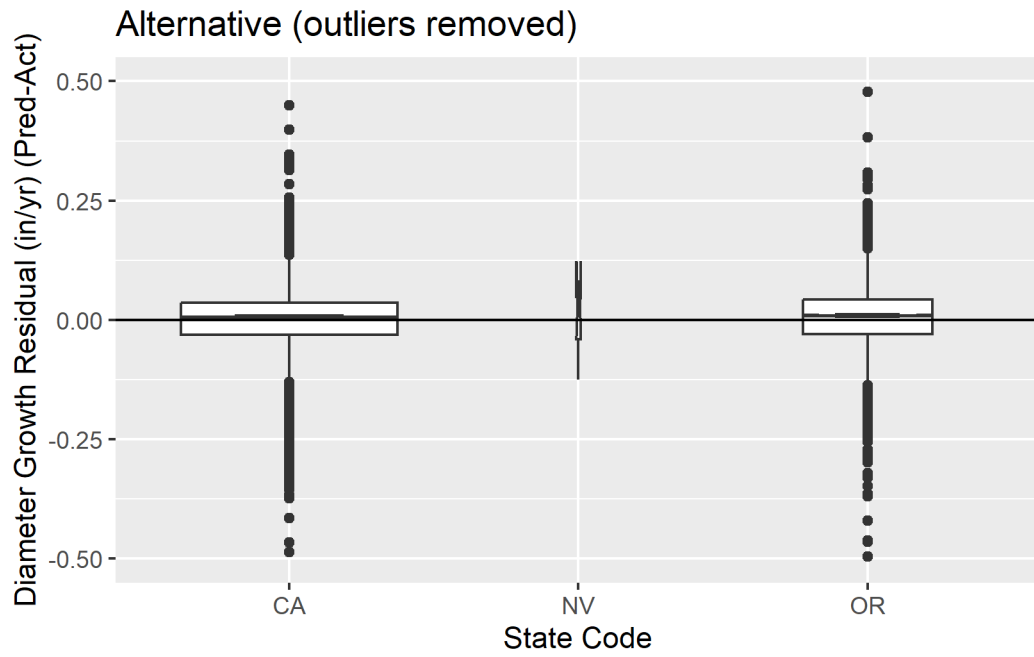


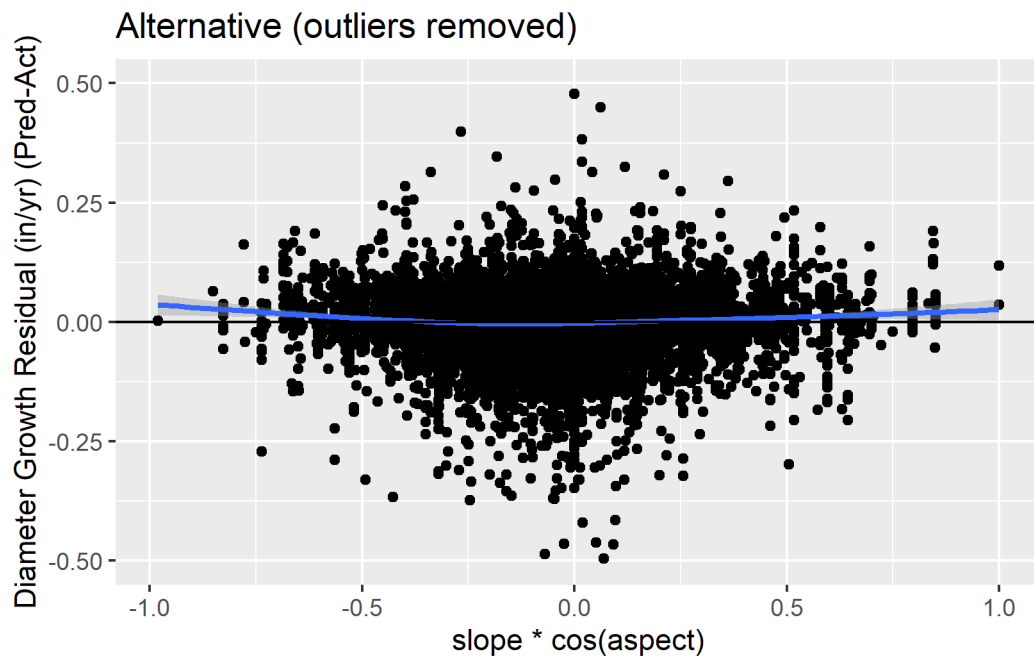
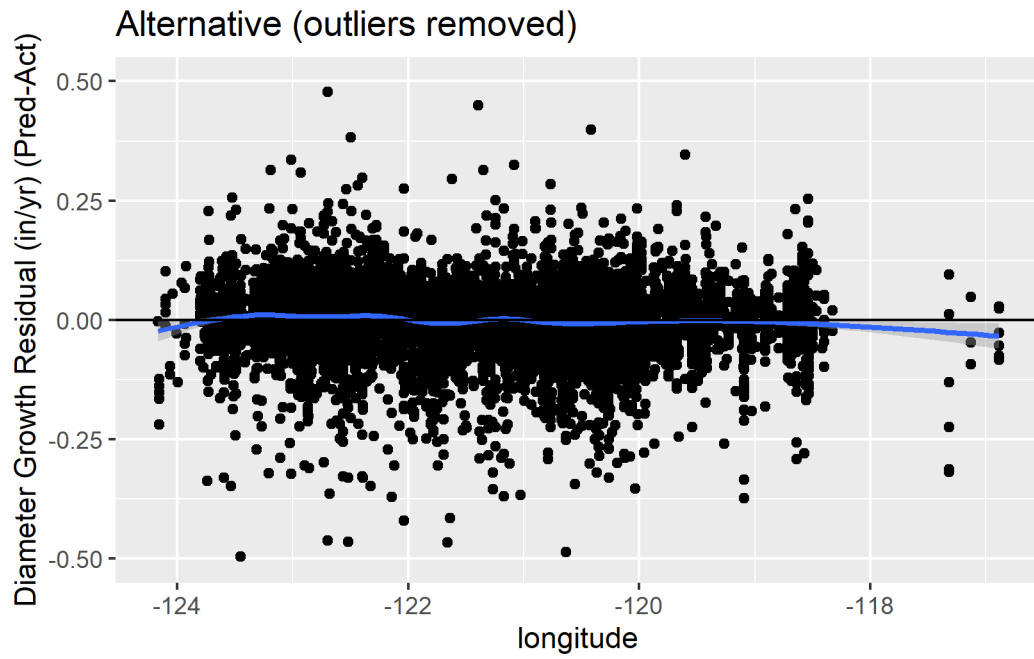
Alternative by si Species (outliers removed)

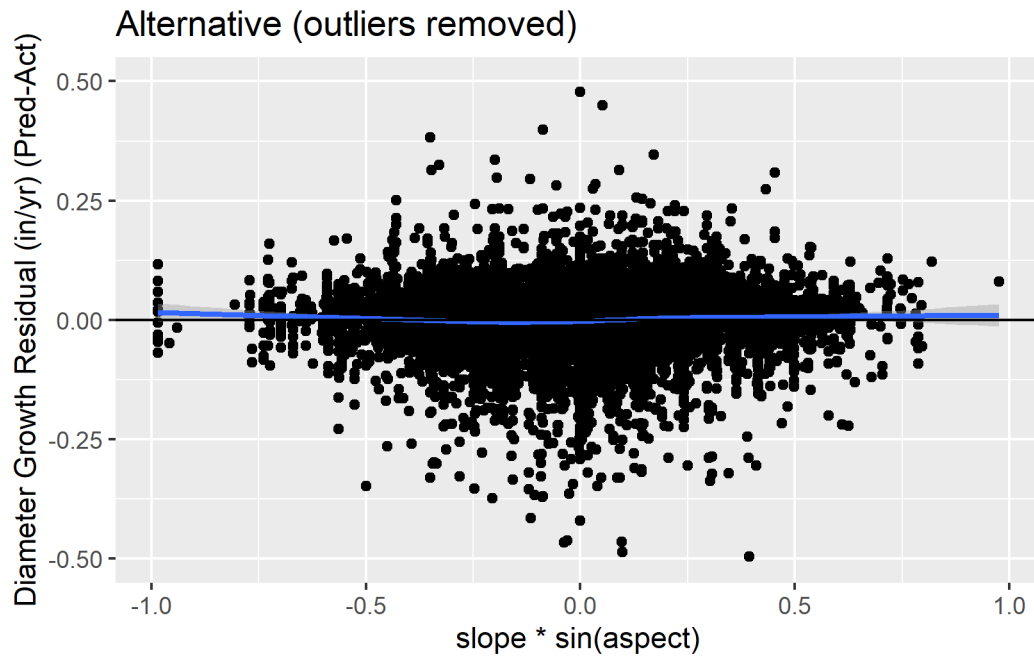


Alternative (outliers removed)

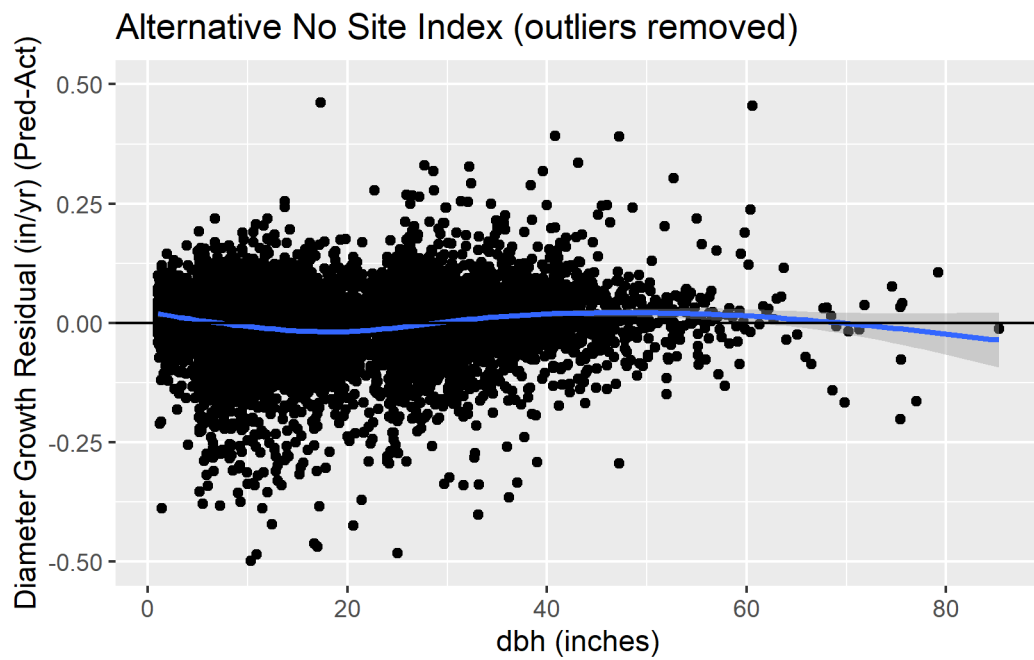


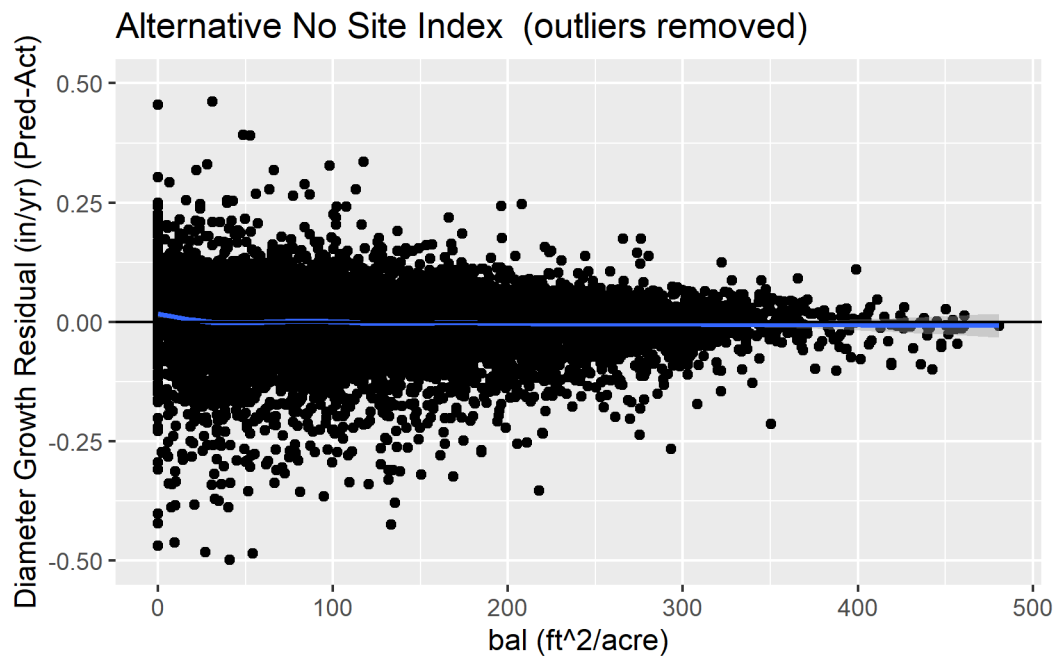
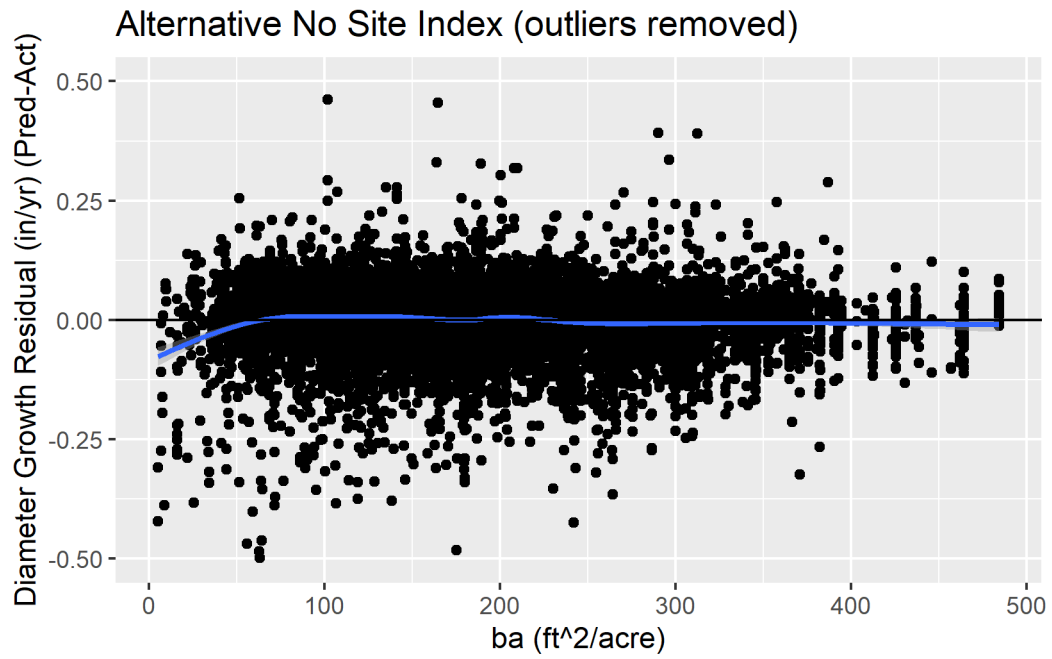


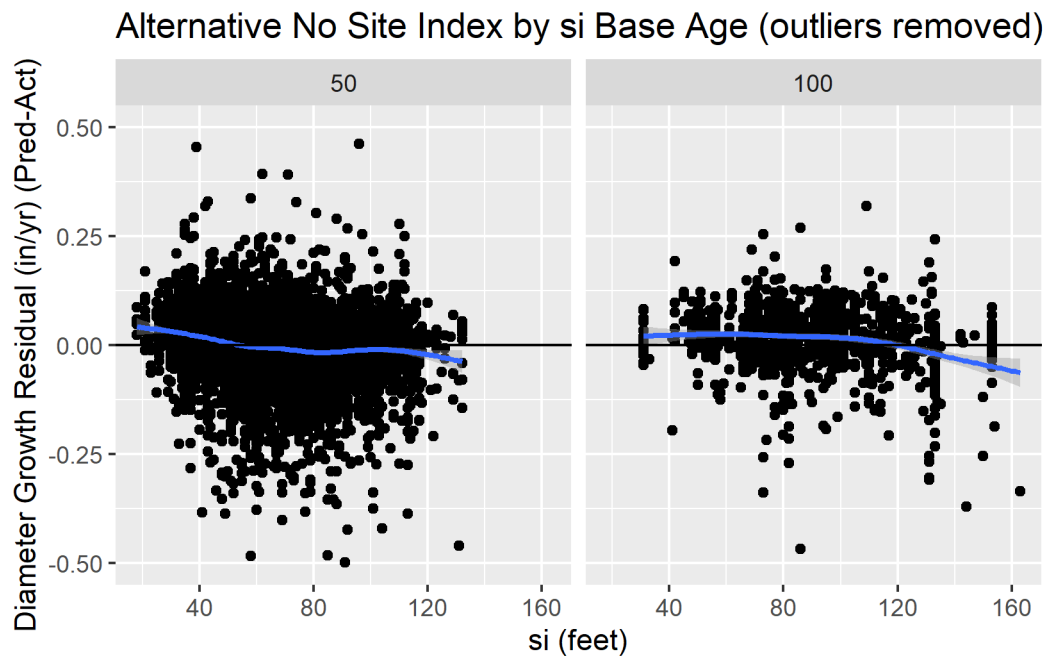
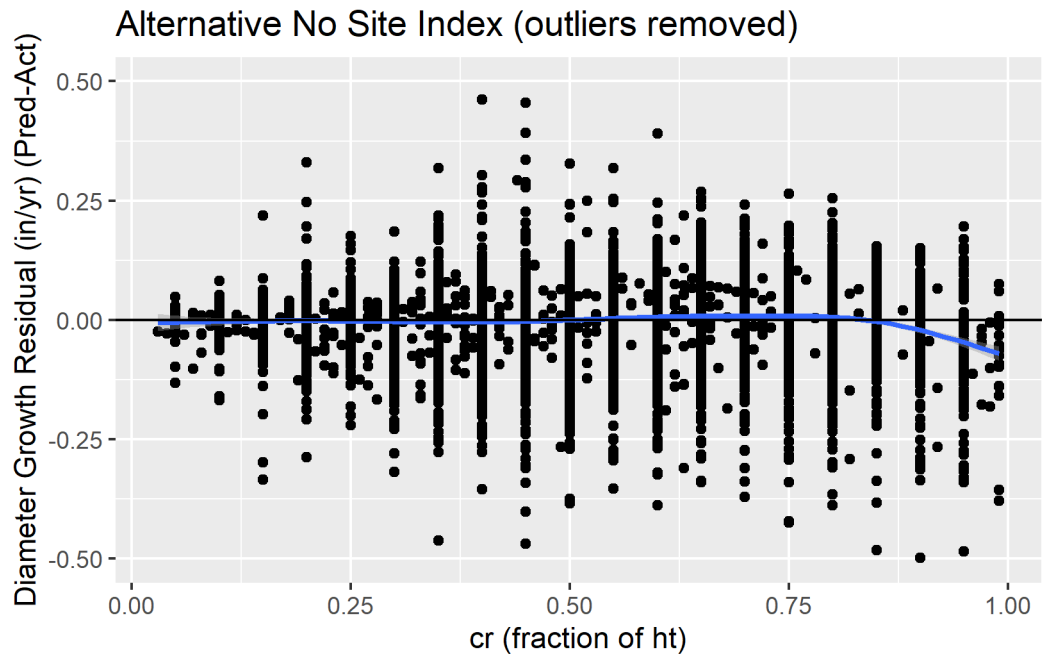




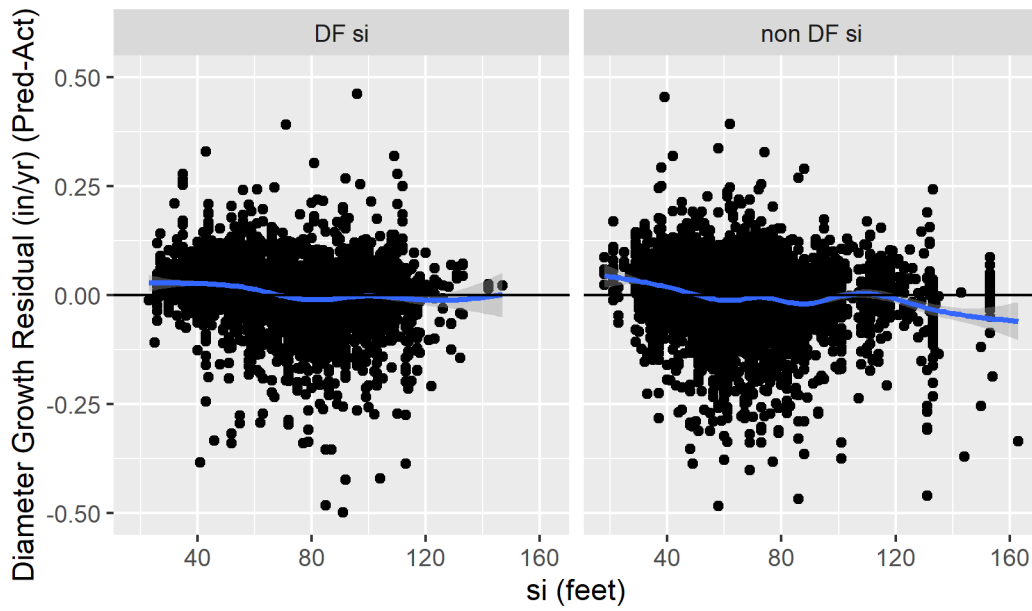
Residual Analysis for Equation 2



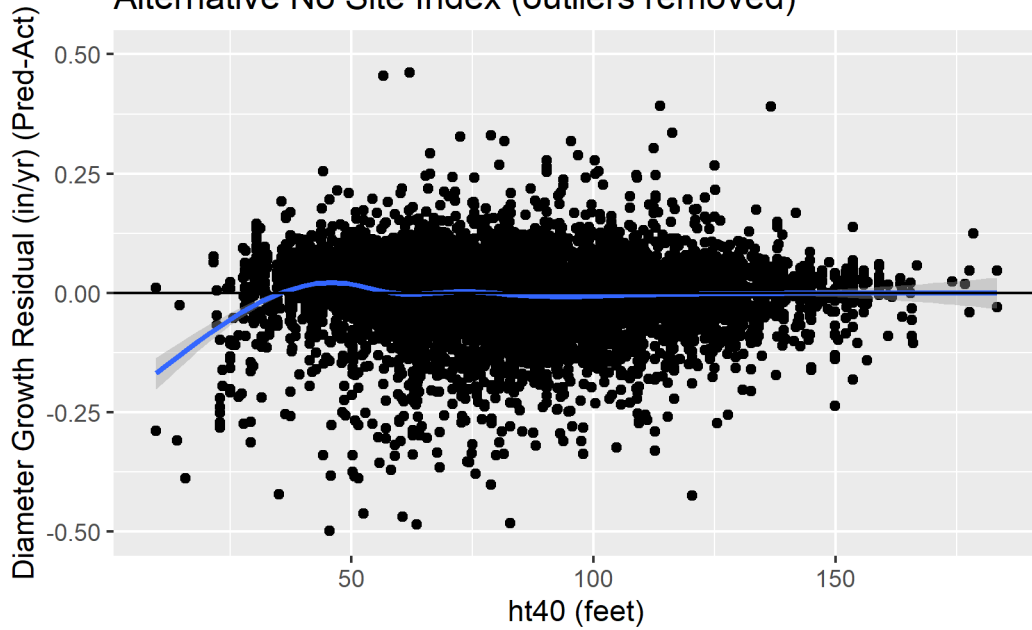


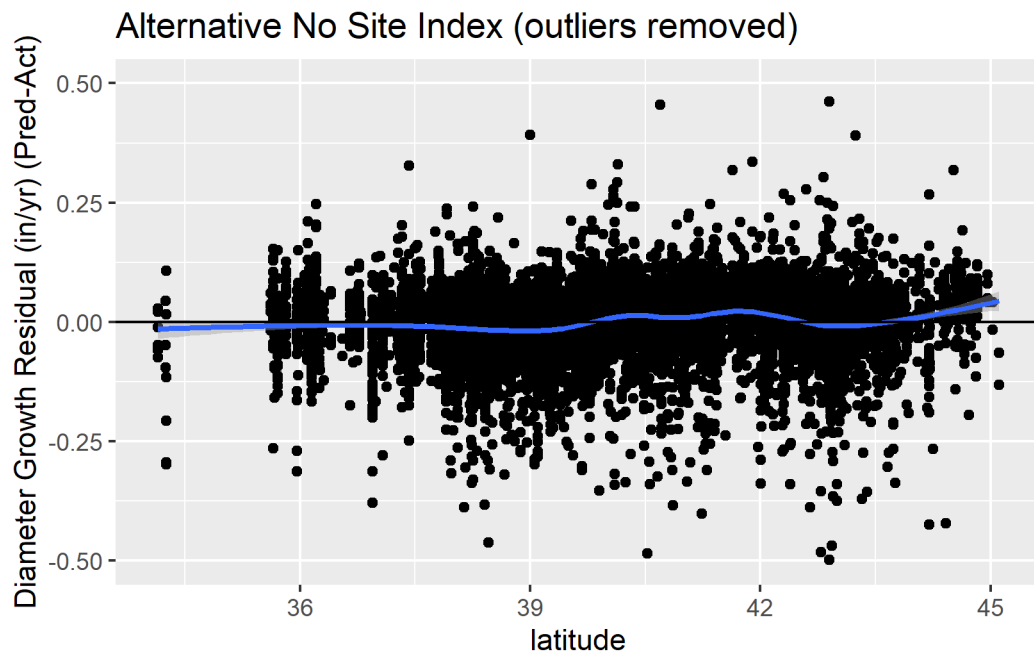
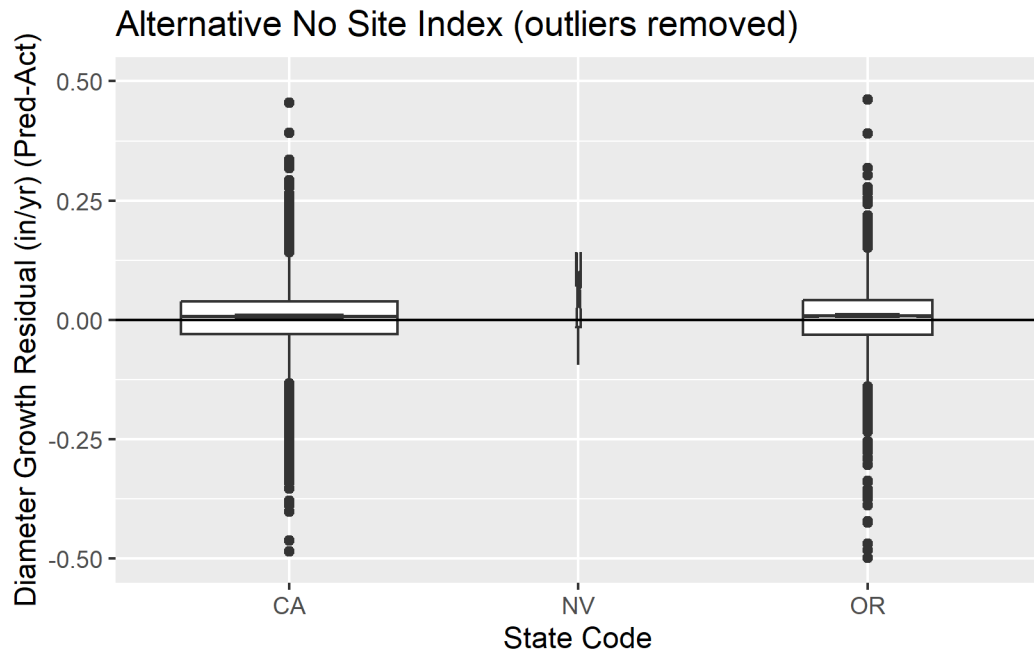


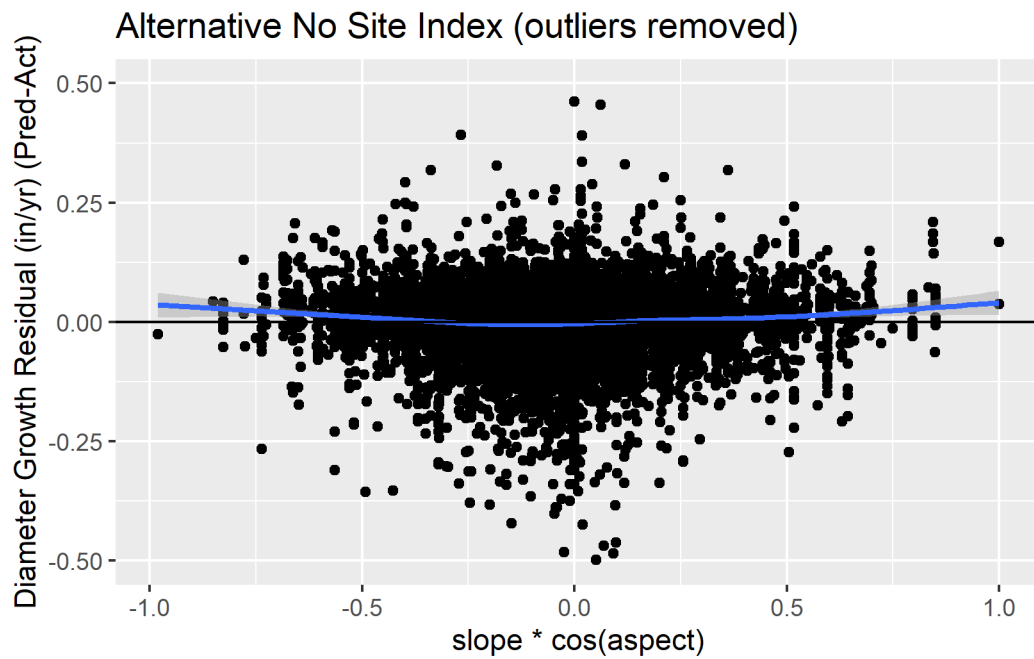
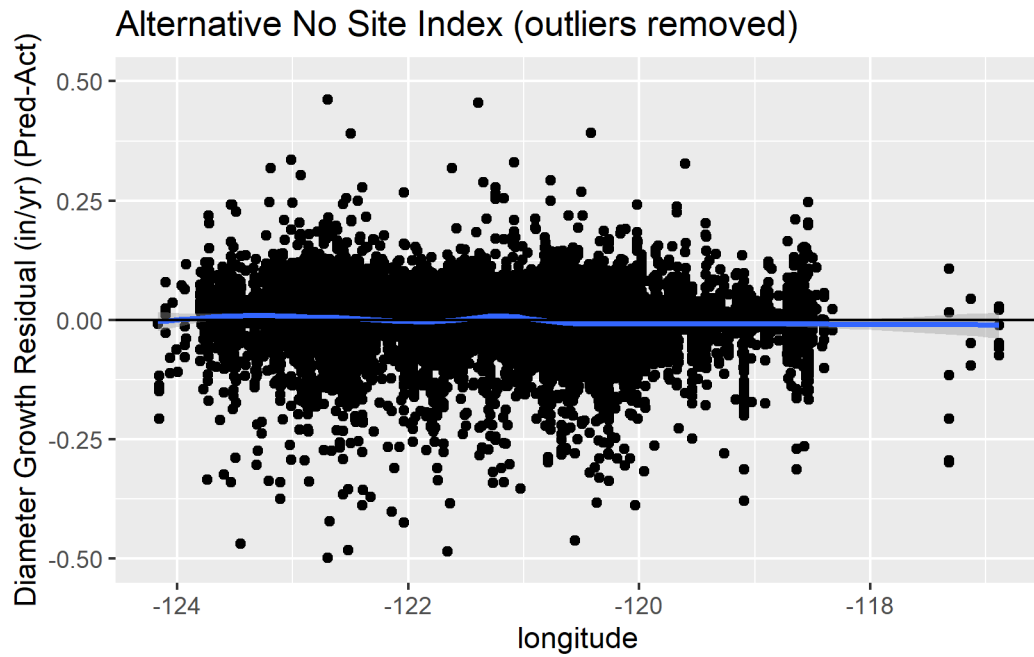
Alternative No Site Index by si Species (outliers removed)



Alternative No Site Index (outliers removed)







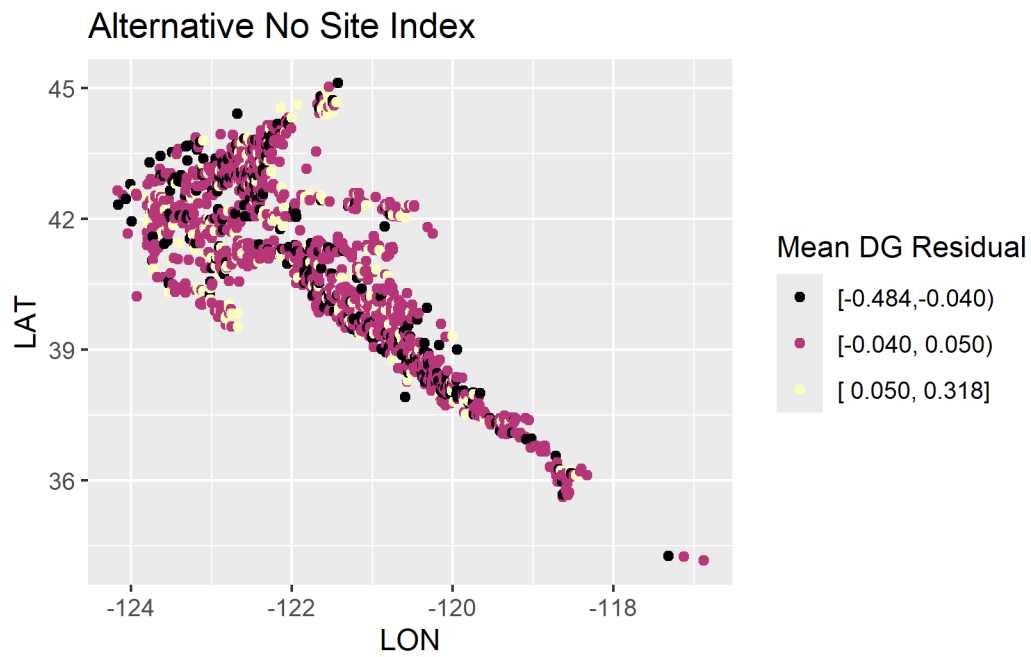
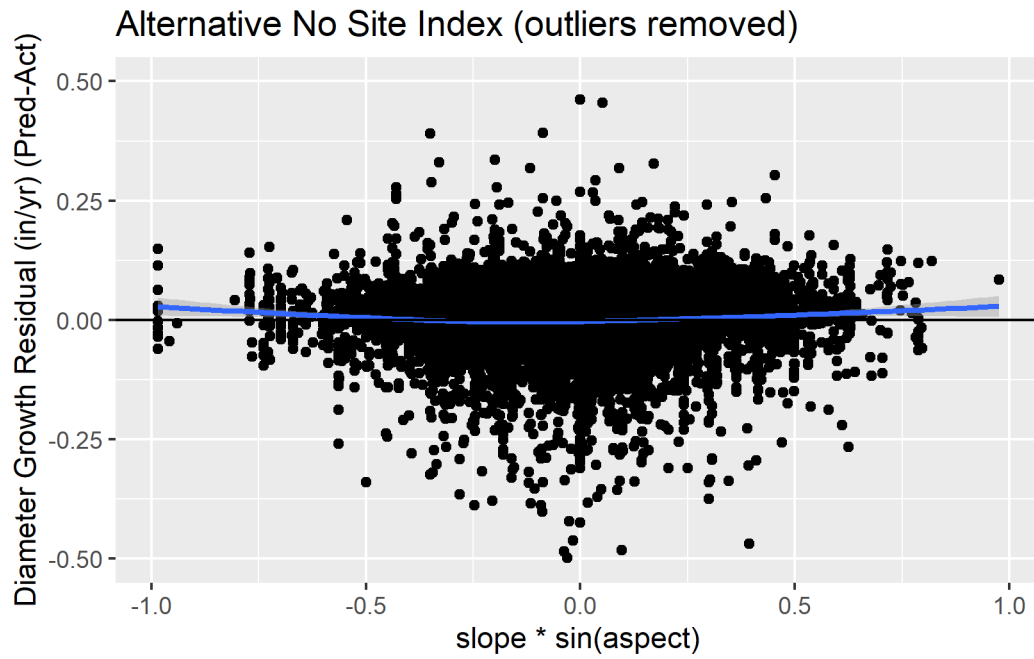


Table 3: Independent Variables for One Inch dbh Trees

Variable	N	Mean	Std. Dev.	Min	Pctl. 25	Pctl. 75	Max
ba	37	179	86	52	117	231	431
bal	37	178	86	52	117	230	430
ht	37	7.3	1.4	5	7	8	12
cr	37	0.42	0.24	0.05	0.2	0.55	0.9
si	37	73	26	30	56	89	129

Discussion

Removing **si** degrades the fit marginally. Otherwise, the alternative model fits the data suprisingly well given the limited data.

Equation Behavior for Very Small Trees

