

Alternative Red Alder Diameter Growth

Greg Johnson and David Marshall

2026-01-20

Data

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

After subsetting the data to censor observations with missing data, limiting the species to Red Alder (FIA species code 351), and remeasurement intervals ≥ 5 years we get the observations in Table 1.

Table 1: Red Alder Growth Observations by State

State	Observations
CA	518
OR	4087
WA	4538

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 Red Alder 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 18 different site index equations for 16 species. Red Alder site index comprises 11% 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 (*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.

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
```

27352.17 27409.14 -13668.09

Random effects:

Formula: B3 ~ 1 | SIINT

B3 Residual

StdDev: 0.03030948 1.078242

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

	Value	Std.Error	DF	t-value	p-value
B0	-2.4626869	0.25910501	9134	-9.50459	0e+00
B1	-0.7725991	0.02002707	9134	-38.57775	0e+00
B2	-0.1760181	0.04622253	9134	-3.80806	1e-04
B3	0.3477299	0.05071774	9134	6.85618	0e+00
B4	1.2842251	0.03460138	9134	37.11485	0e+00
B5	0.5334722	0.04360240	9134	12.23493	0e+00

Correlation:

	B0	B1	B2	B3	B4
B1	-0.211				
B2	-0.387	0.489			
B3	-0.824	-0.022	0.013		
B4	-0.420	0.312	0.510	-0.038	
B5	-0.371	0.471	0.994	0.018	0.472

Standardized Within-Group Residuals:

	Min	Q1	Med	Q3	Max
	-48.0077227	-0.4987108	-0.1071605	0.4341391	8.7272421

Number of Observations: 9143

Number of Groups: 4

\$SIINT

	B3
50.FALSE	0.025303939
100.FALSE	-0.000699833
50.TRUE	0.020081107
100.TRUE	-0.044685213

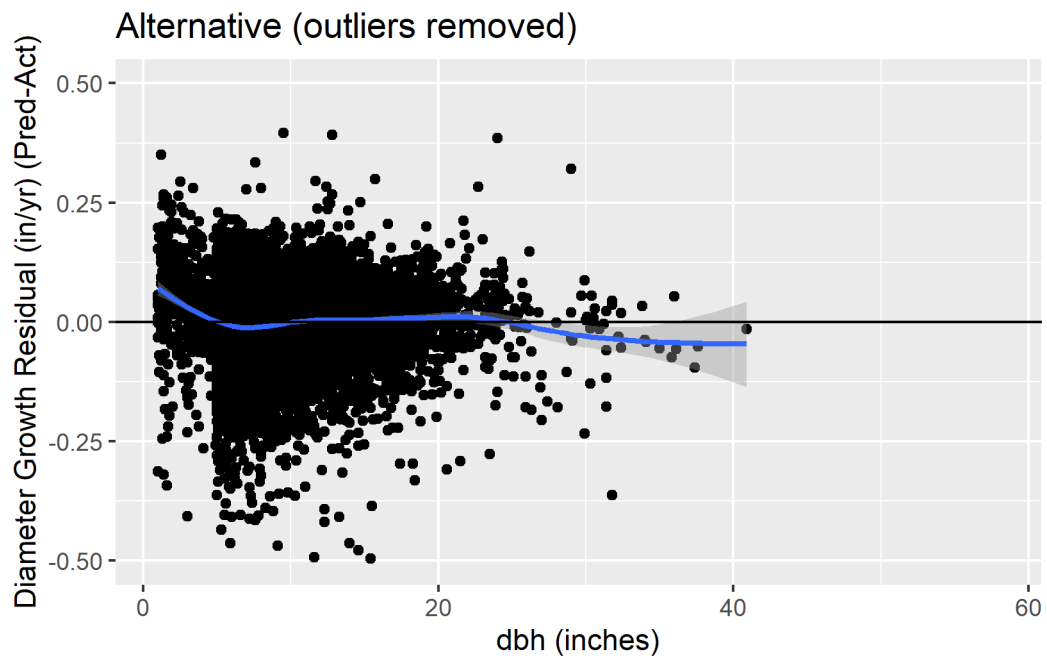
Residual Standard Error: 1.07824246806703 on 9134 degrees of freedom, AIC: 27352.2

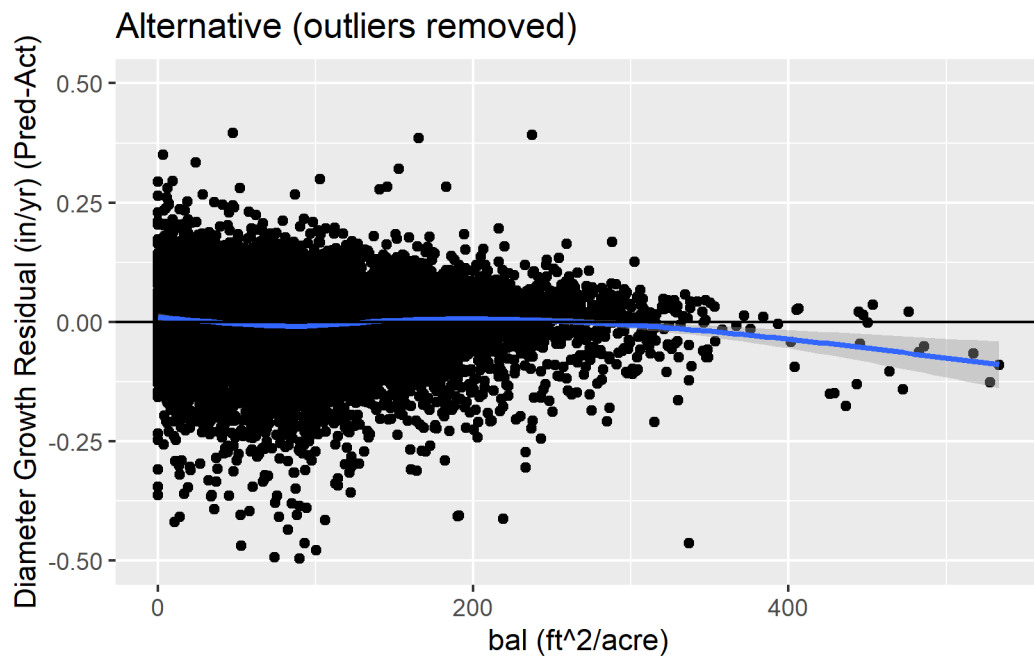
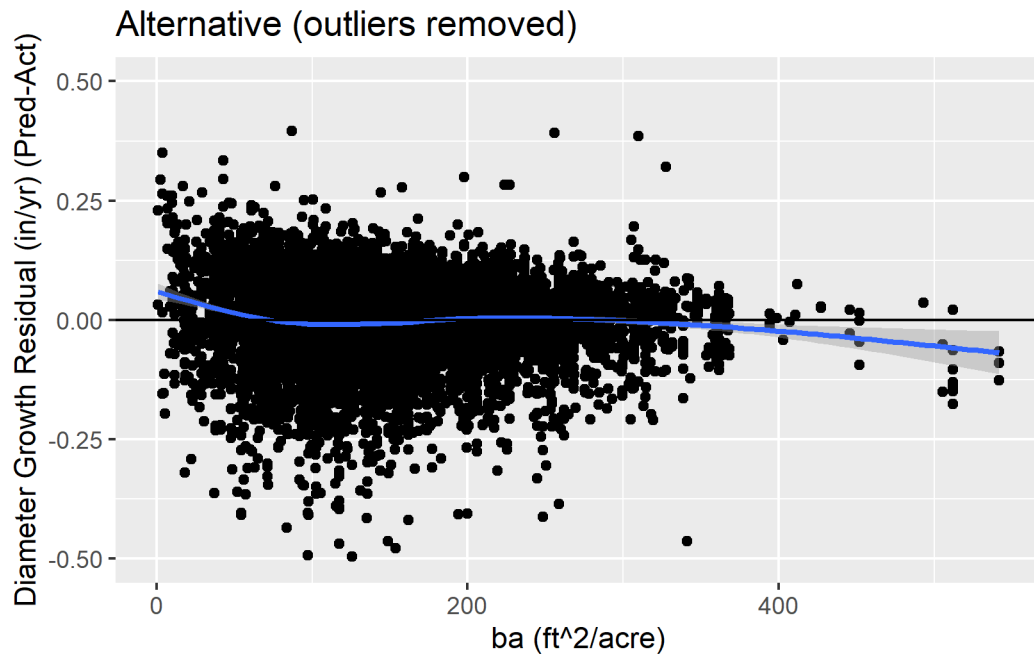
and for Equation 2:

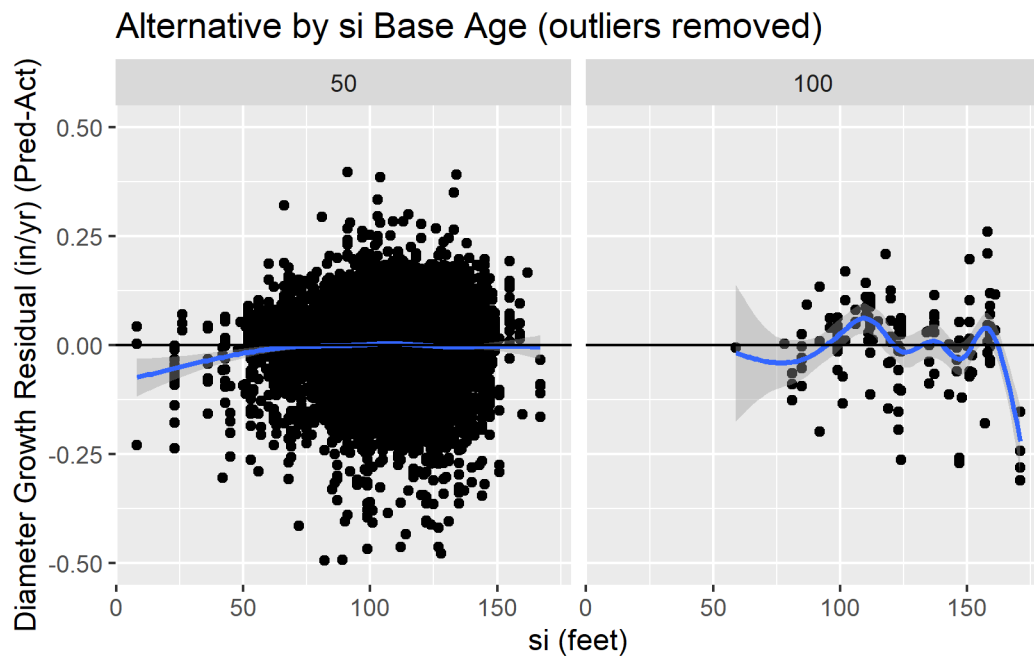
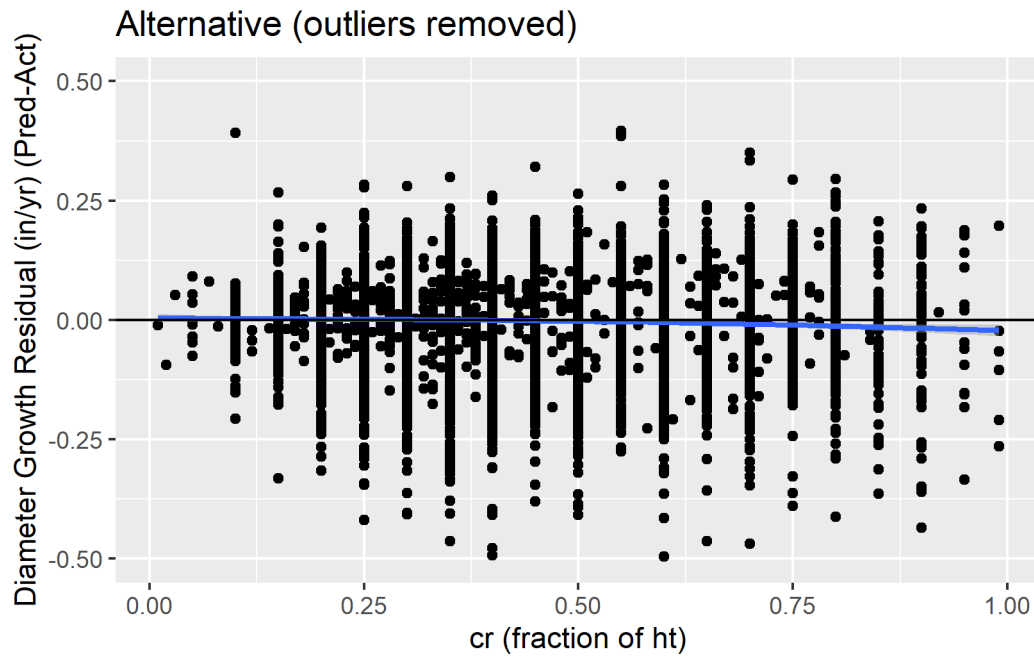
	Coef.	Std. error	t-stat.	p
B0	-0.7667739	0.1302019	-5.889116	0.0000000
B1	-0.7739417	0.0201284	-38.450326	0.0000000
B2	-0.1836459	0.0486083	-3.778073	0.0001591
B4	1.3021968	0.0349787	37.228288	0.0000000
B5	0.5232333	0.0437591	11.957122	0.0000000

Residual Standard Error: 1.08259291867095 on 9138 degrees of freedom, AIC: 27404.9

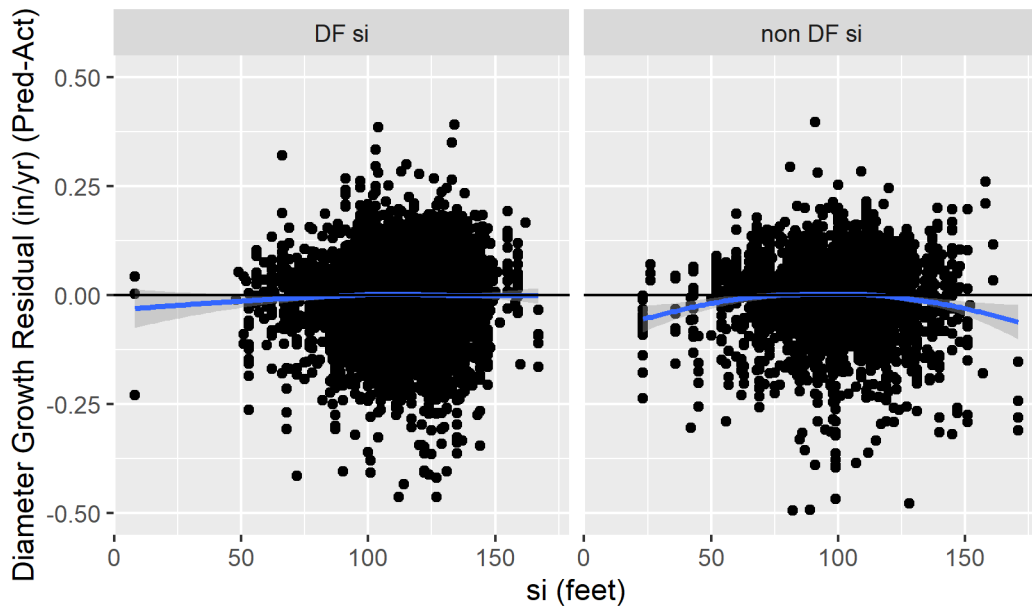
Residual Analysis for Equation 1



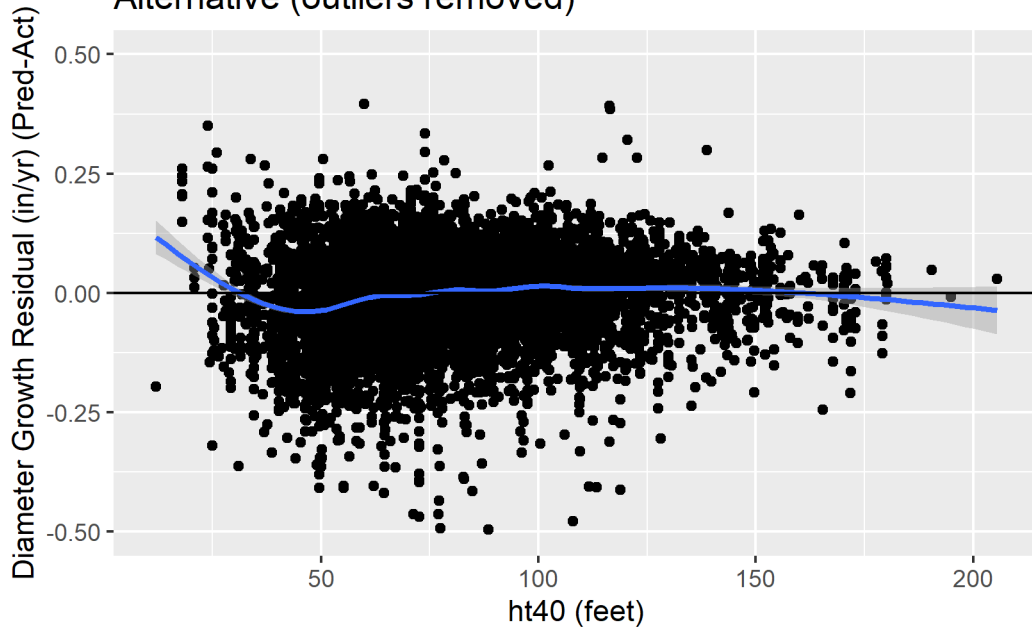


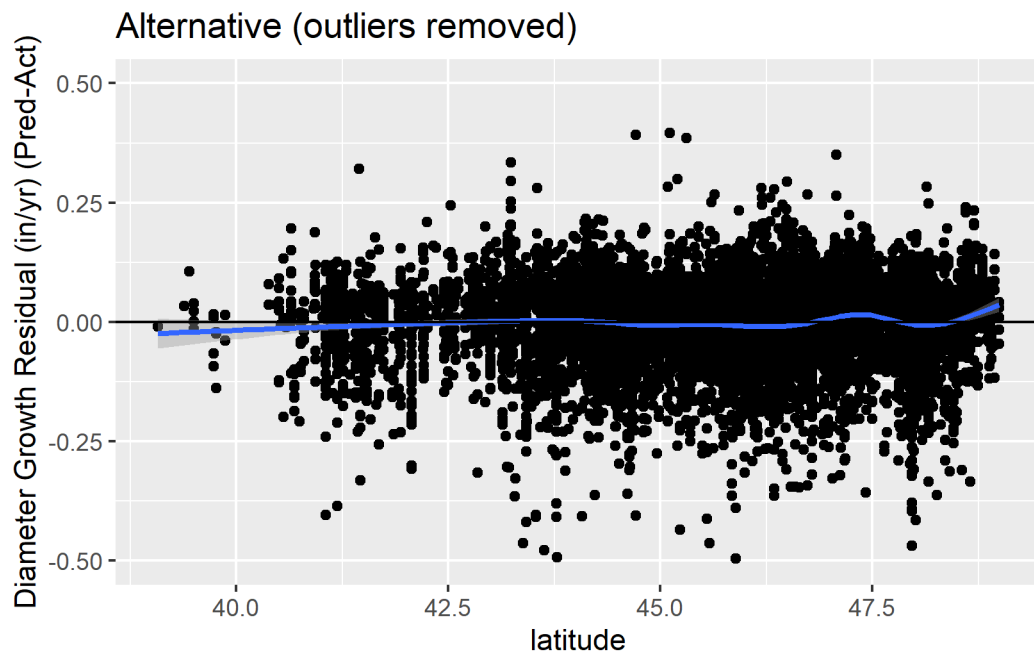
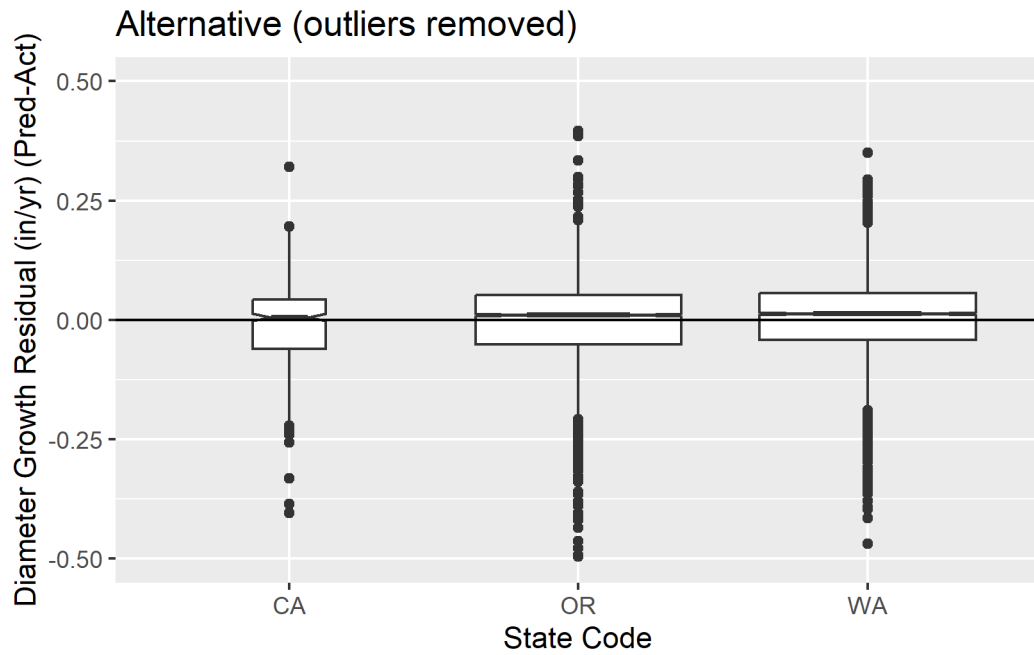


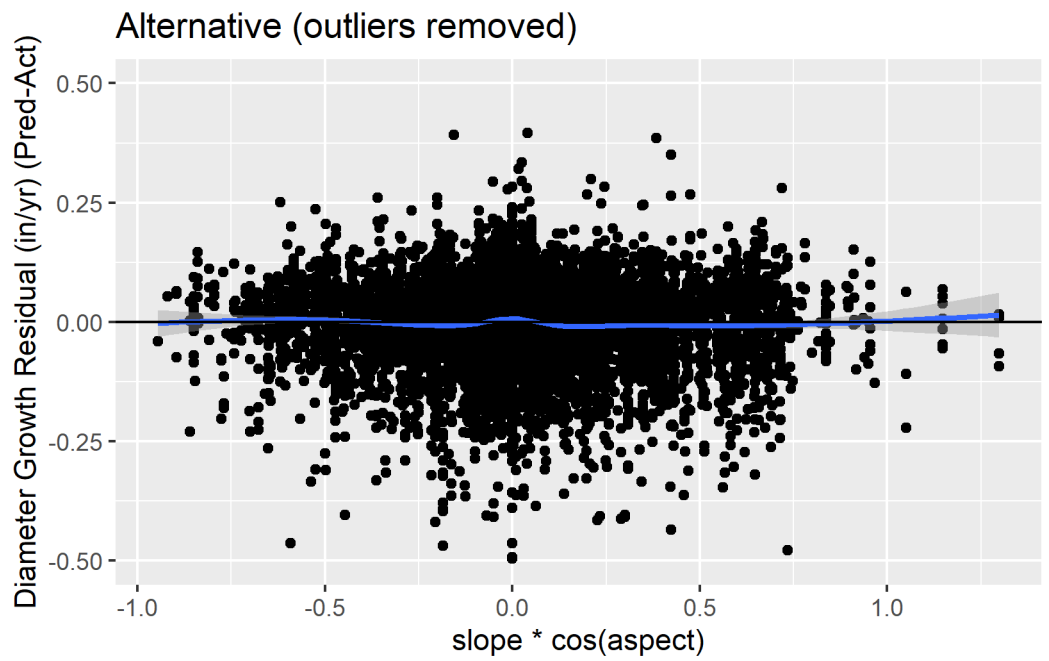
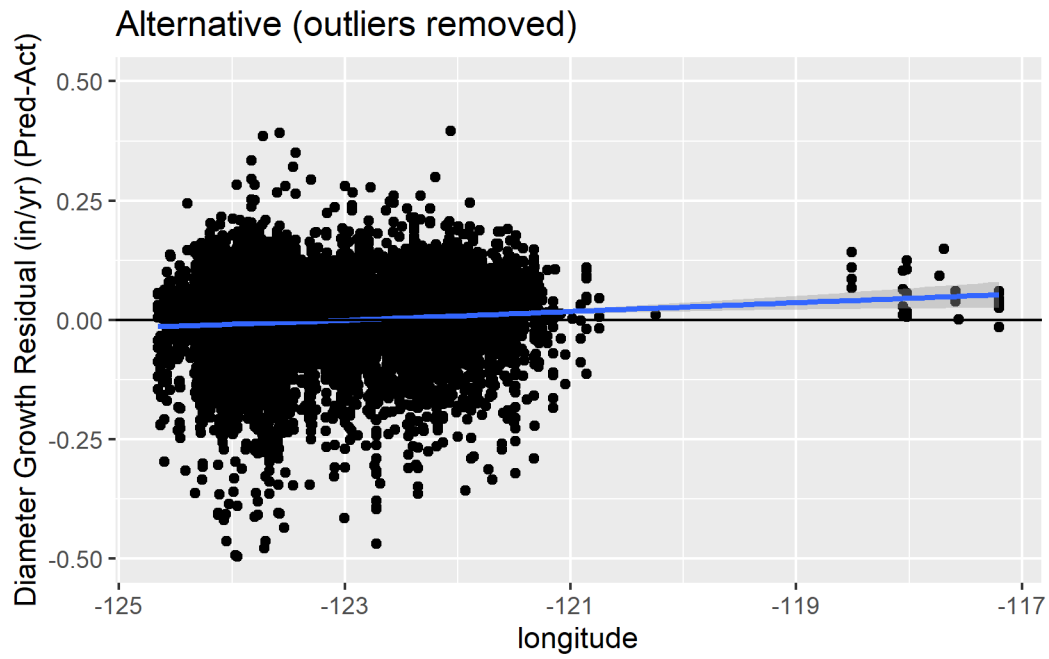
Alternative by si Species (outliers removed)

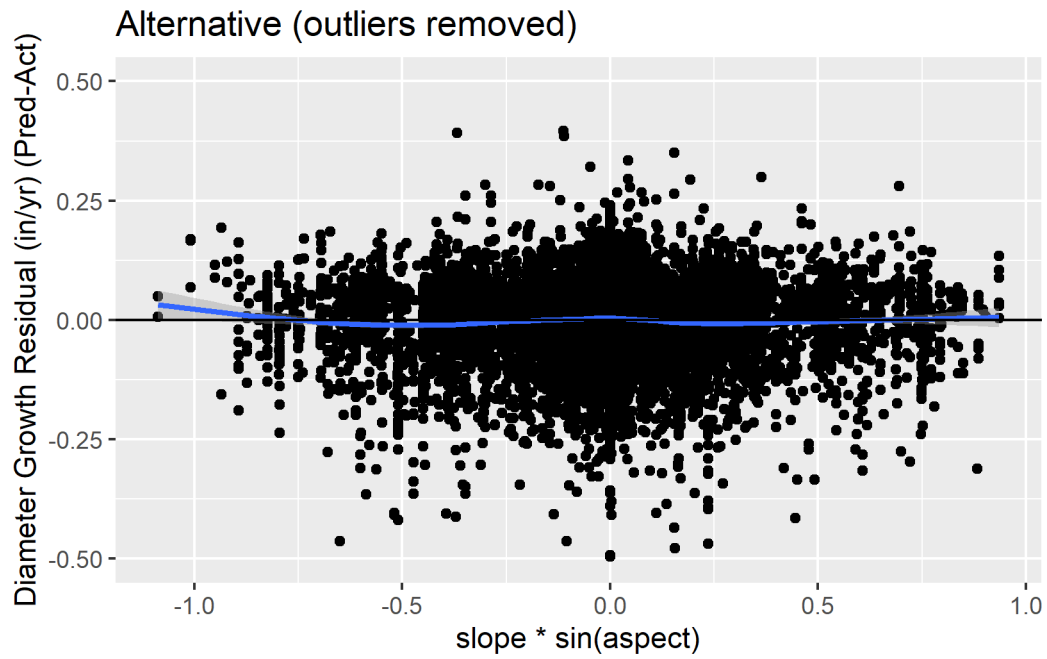


Alternative (outliers removed)

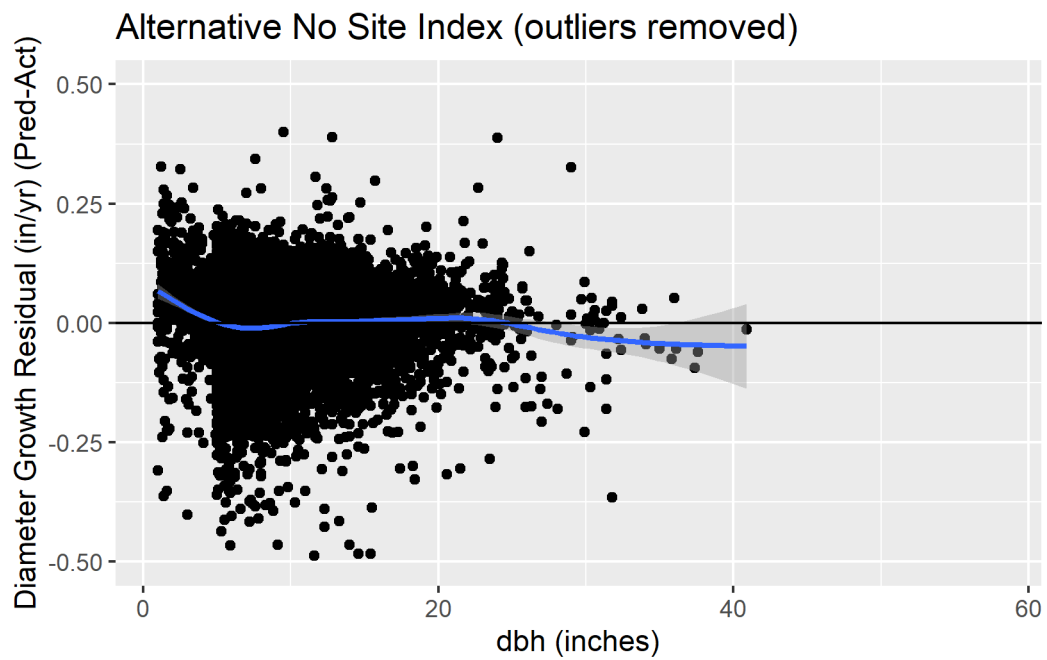


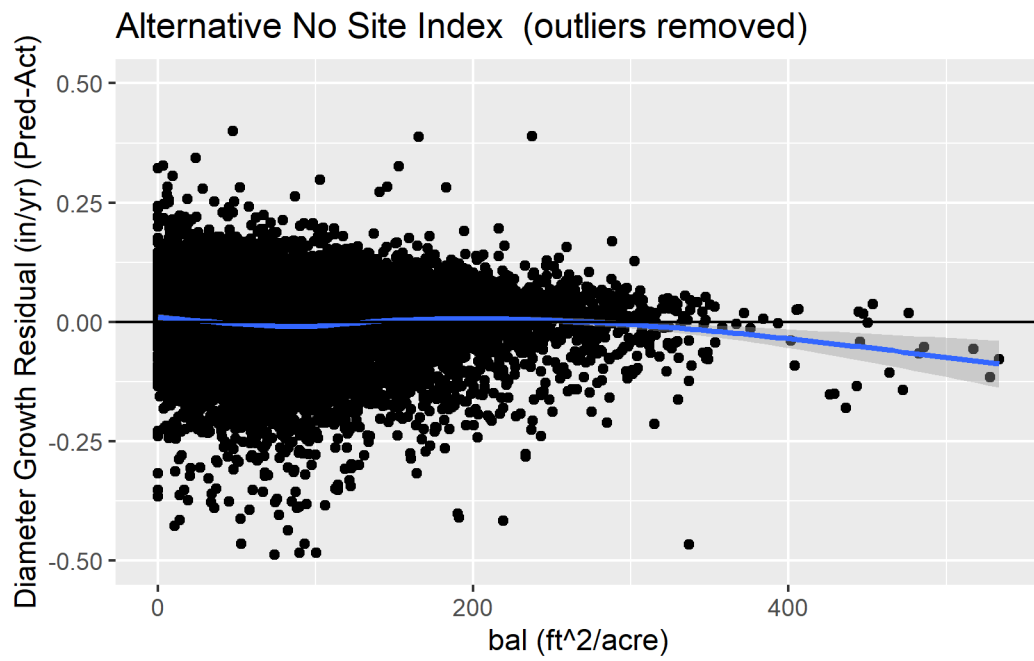
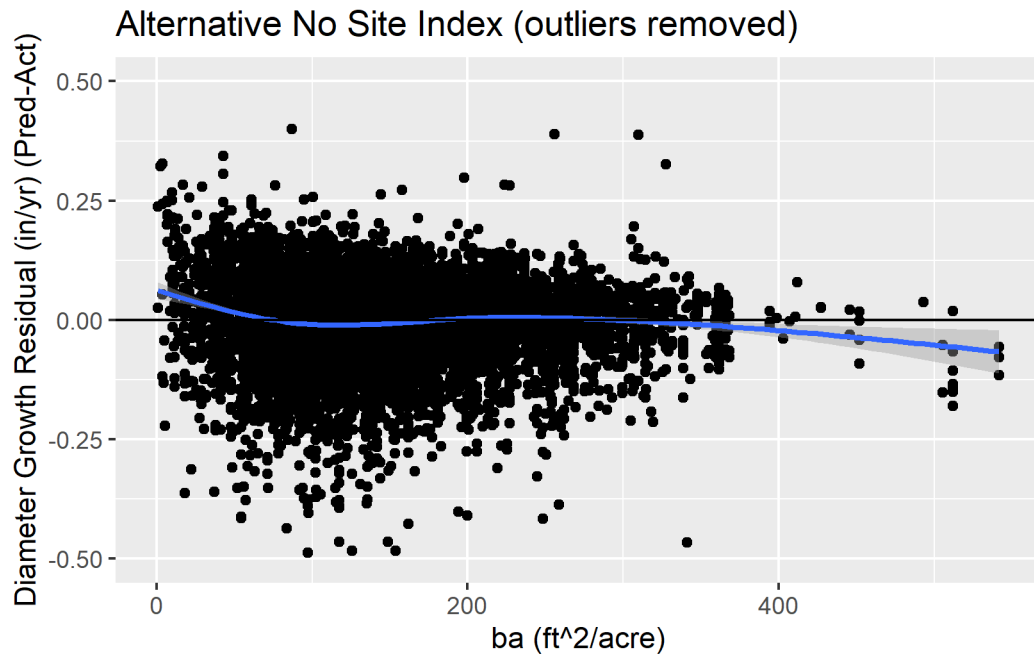


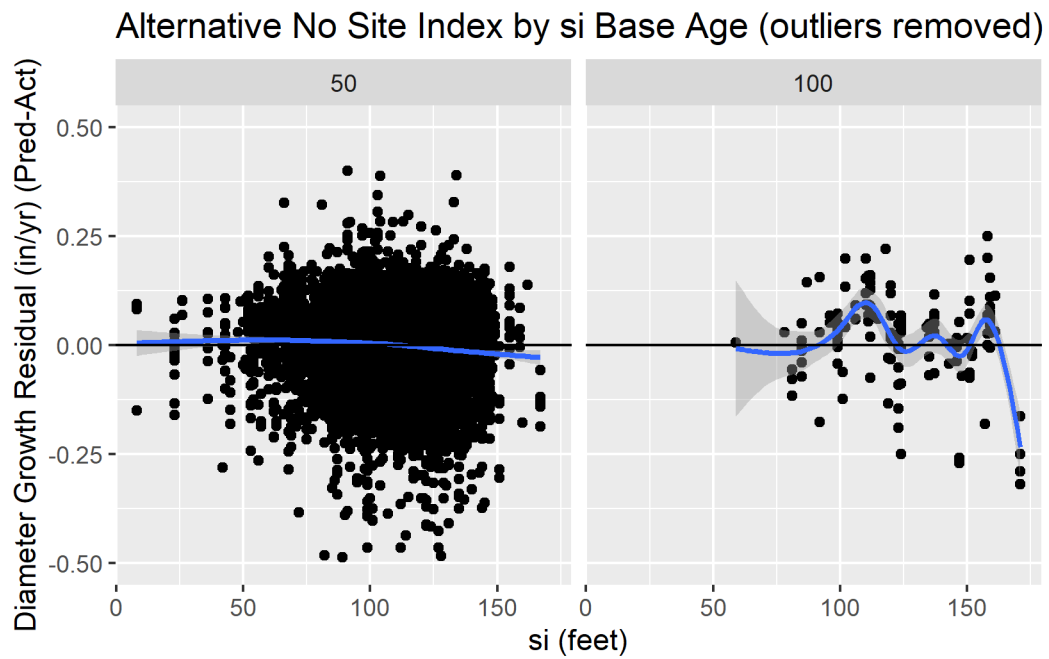
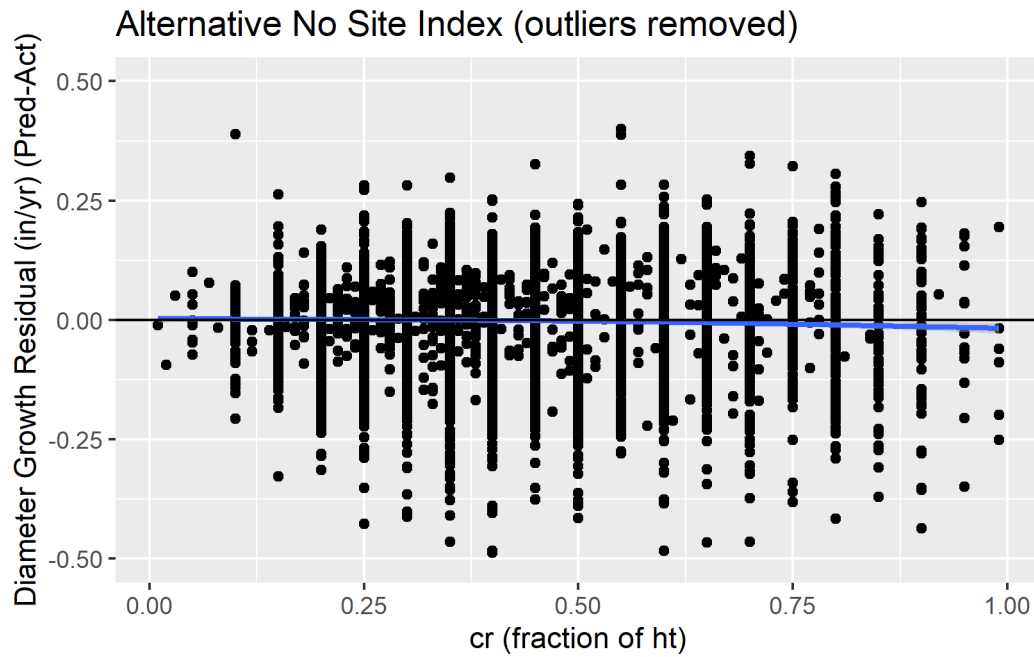




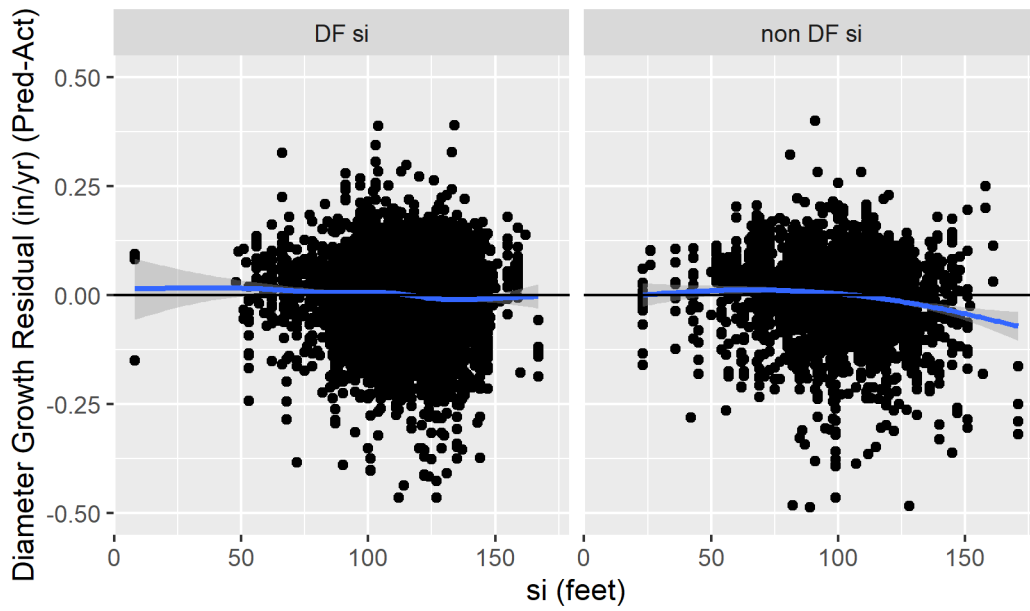
Residual Analysis for Equation 2



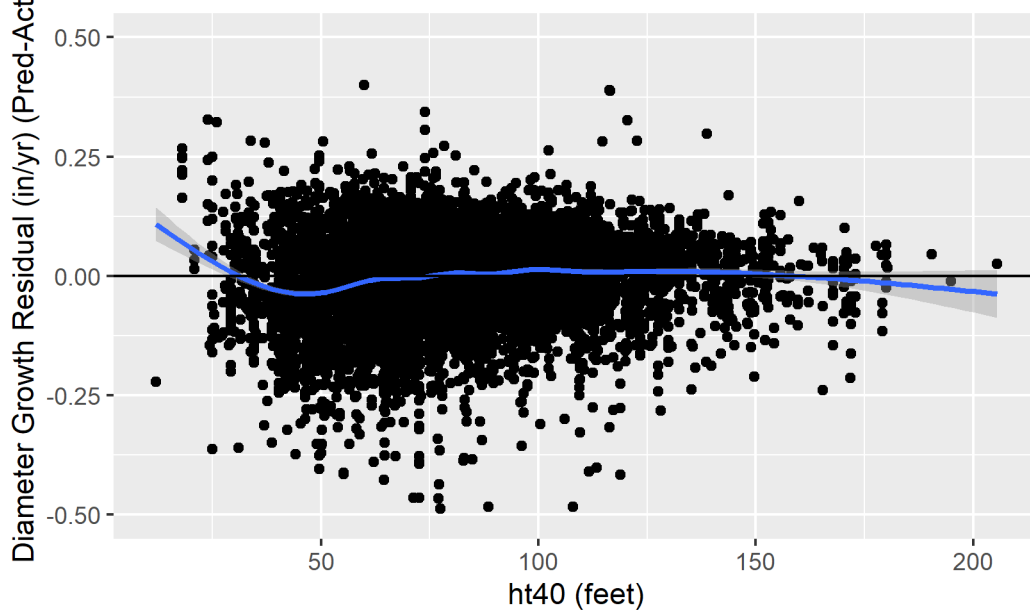


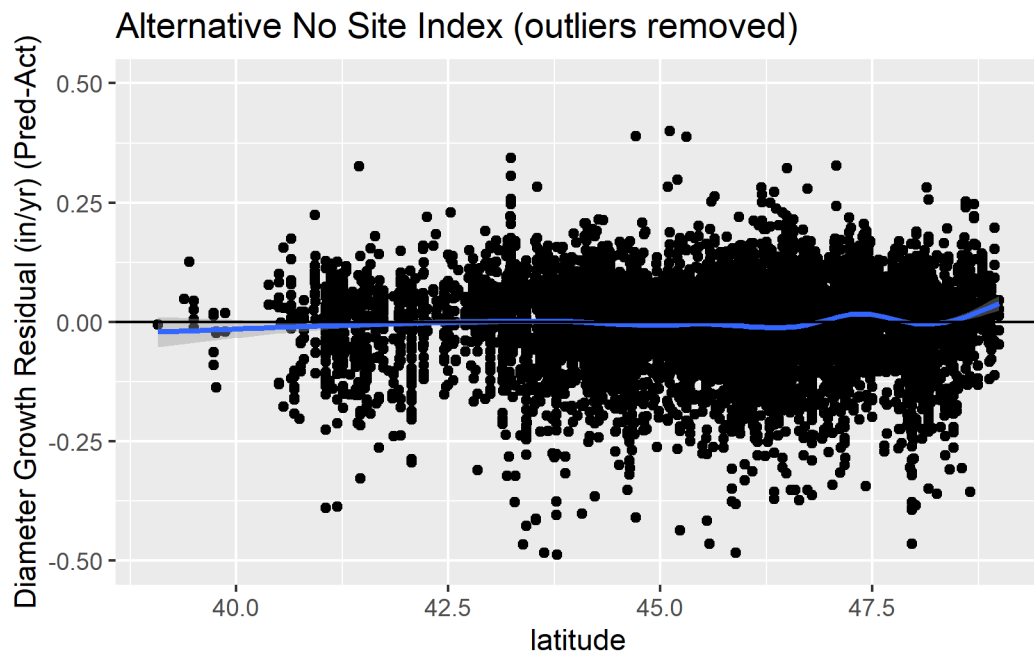
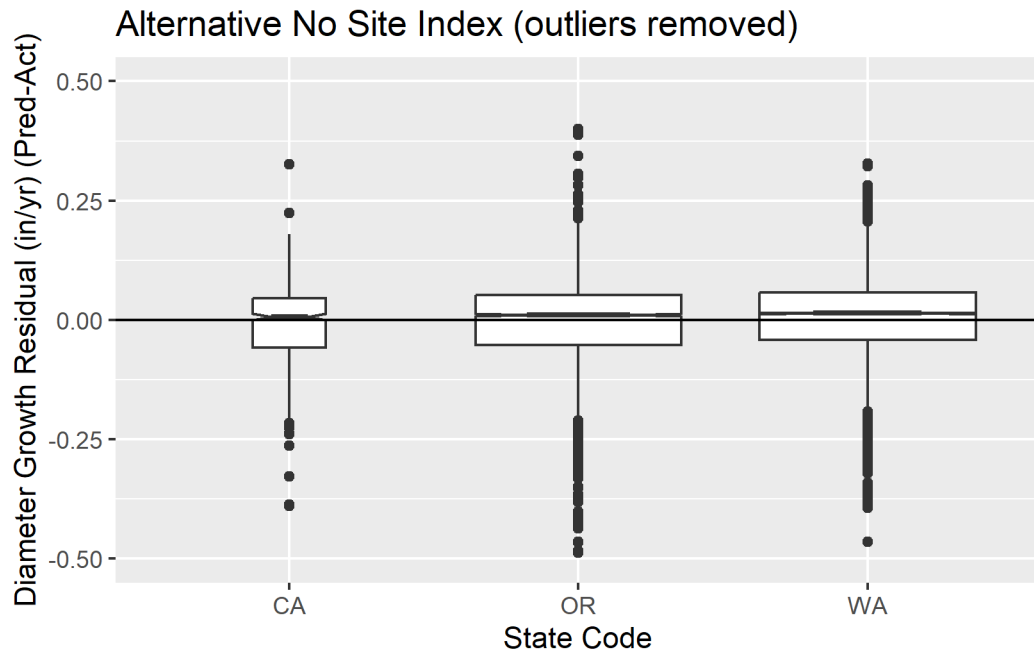


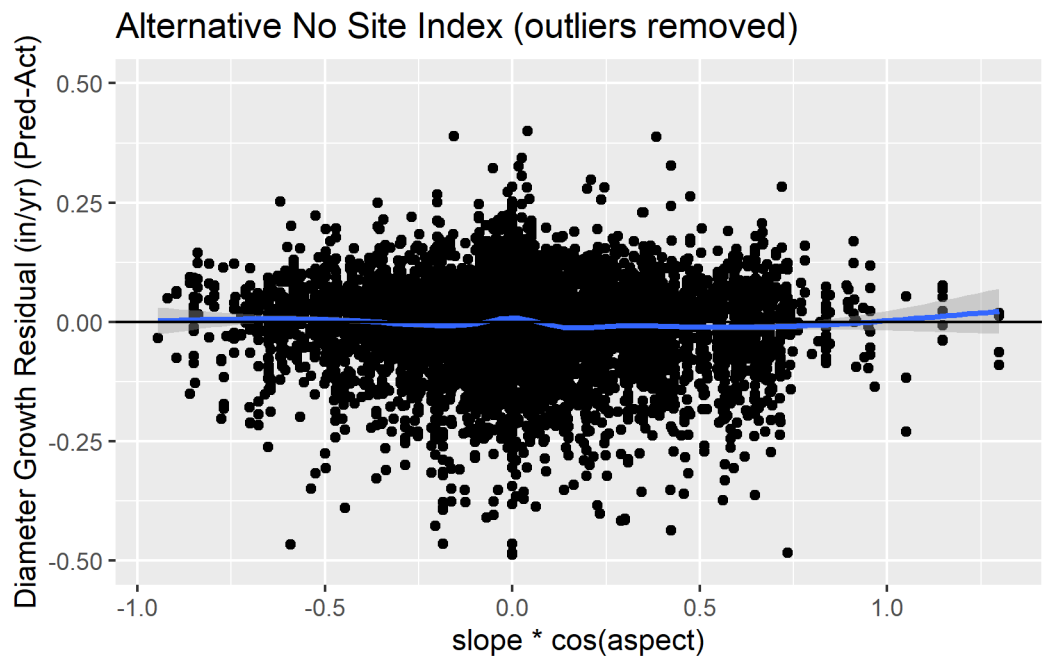
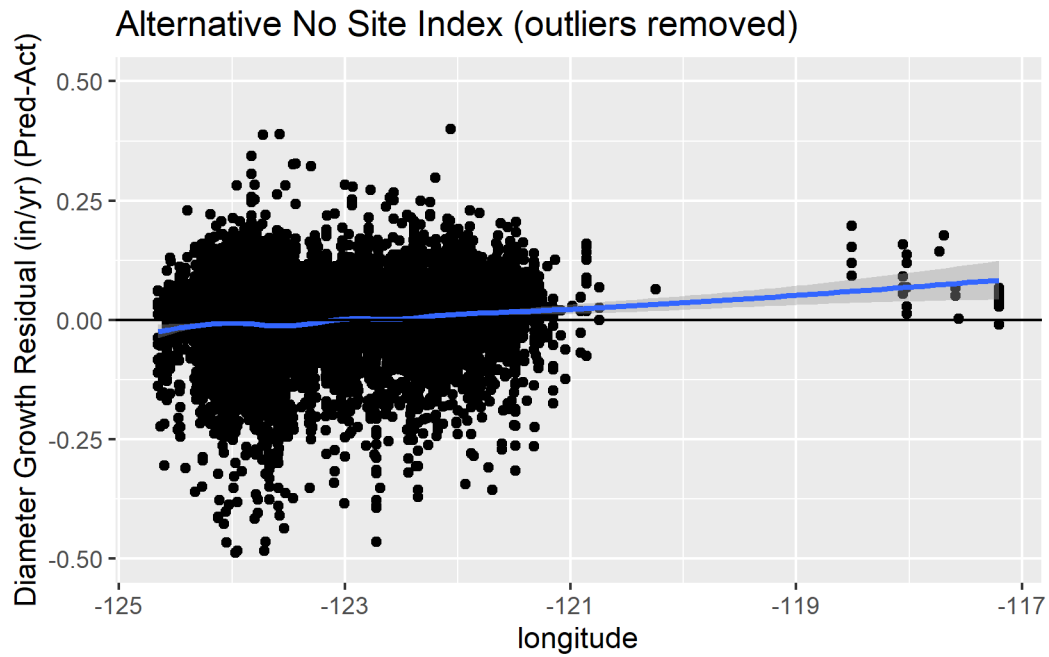
Alternative No Site Index by si Species (outliers removed)



Alternative No Site Index (outliers removed)







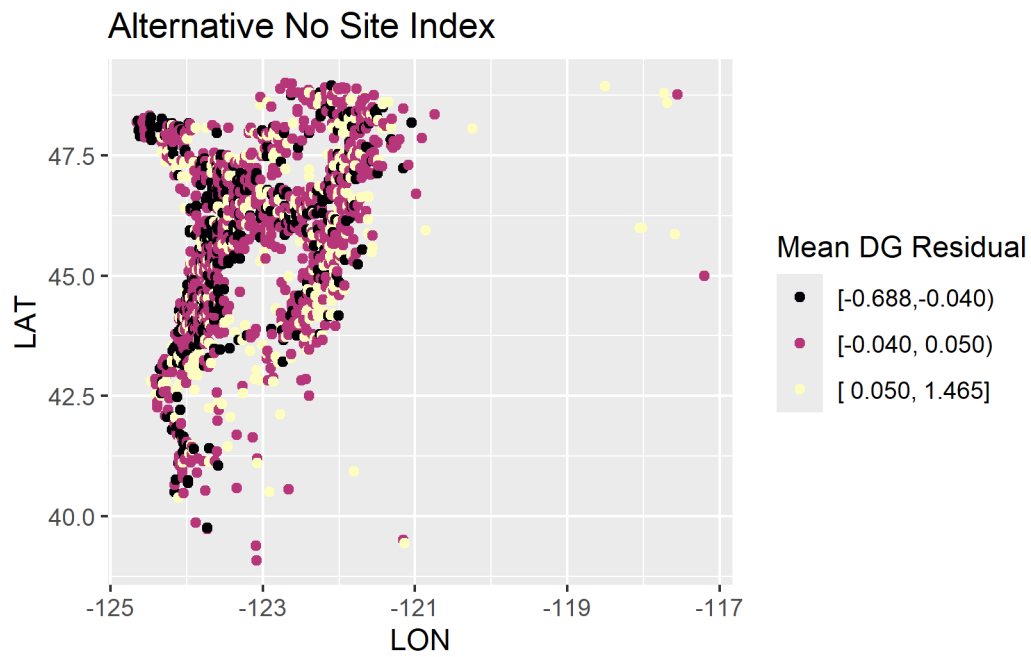
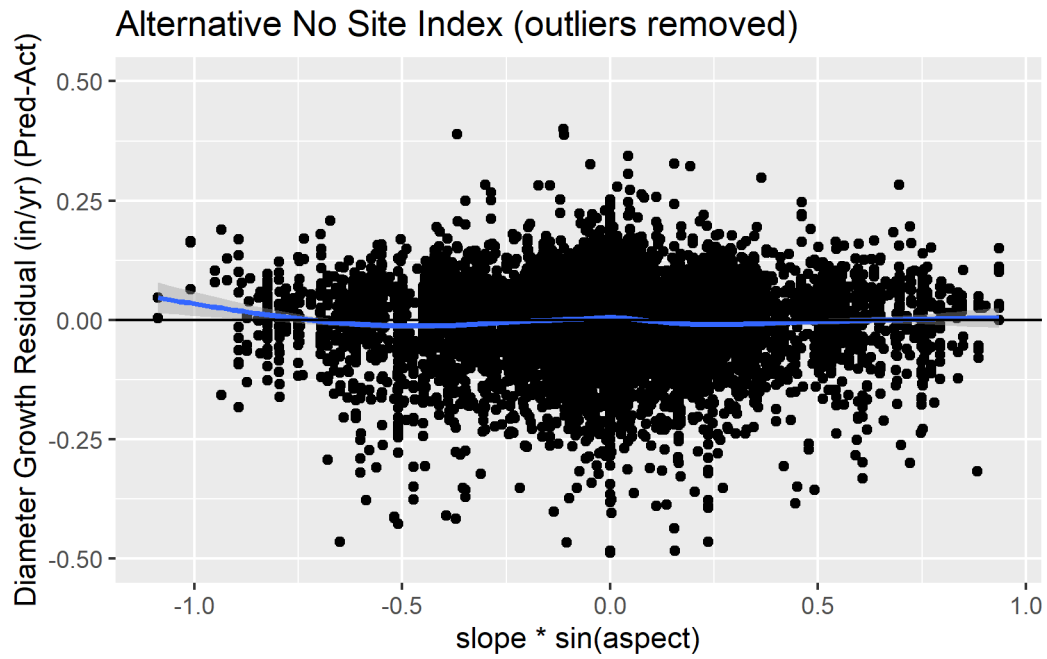


Table 3: Independent Variables for One Inch dbh Trees

Variable	N	Mean	Std. Dev.	Min	Pctl. 25	Pctl. 75	Max
ba	11	101	74	15	57	109	238
bal	11	100	74	15	57	108	237
ht	11	13	2.2	9	12	14	17
cr	11	0.49	0.26	0.2	0.3	0.6	0.99
si	11	123	18	97	114	134	151

Discussion

Removing **si** degrades the fit marginally. There seems to be an issue with red alder with small **dbh** values that will need to be investigated. the effect of this over-prediction is evident in the behavior of small trees shown in the next section. It is entirely possible that the limited data available in small trees is affecting the equation's performance.

Equation Behavior for Very Small Trees

