

# Alternative Western Hemlock Diameter Growth

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## Data

We extracted and processed Forest Inventory and Analysis (FIA) data from 6 states listed in the native range of Western Hemlock in the Silvics of North America.<sup>1</sup>

After subsetting the data to censor observations with missing data, limiting the species to Western Hemlock (FIA species code 263), and remeasurement intervals  $\geq 5$  years we get the observations in Table 1.

Table 1: Western Hemlock Growth Observations by State

State	Observations
AK	11231
CA	164
ID	1538
MT	561
OR	13035
WA	26360

## Alternative Model Formulation

An alternative to the ORGANON diameter growth equation<sup>2</sup> which reduces parameter count while retaining key features of the original model is shown below. The key change is the

<sup>1</sup>Burns, 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.

<sup>2</sup>Hann, 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.

term with a ratio of a transformation of diameter at breast height ( $\text{dbh}$ ) squared to crown length. Since  $\beta_1$  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 ( $\text{bal}$ ) serves as the inter-tree competition factor, and site index ( $\text{si}$ ) 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 Western Hemlock site trees,
2. It is estimated using a number of different and not necessarily compatible  $\text{si}$  equations, and
3. The available  $\text{si}$  equations do not all use the same base age.

In the data set  $\text{si}$  is derived from 22 different site index equations for 20 species. Western Hemlock site index comprises 45% of the observations. There are 3 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.

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

and

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

where:

- $\text{dbh}$  = diameter at breast height (inches),
- $\text{bal}$  = basal area per acre in larger trees ( $\text{feet}^2/\text{ac}$ ),
- $\text{cr}$  = crown ratio (fraction of total height),
- $\text{ht}$  = total height (feet), and
- $\text{si}_{s,b}$  = 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  $\text{dbh}$ . 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\$SIINT)))  
 AIC          BIC         logLik  
 118654.5  118725.5  -59319.26

Random effects:  
 Formula: B3 ~ 1 | SIINT  
               B3  Residual  
 StdDev: 0.04165923 0.7425849

Fixed effects: B0 + B1 + B2 + B3 + B4 + B5 ~ 1  
 Value  Std.Error    DF    t-value p-value  
 B0 -5.068187 0.06107600 52879 -82.98165 0  
 B1 -0.646654 0.00600721 52879 -107.64641 0  
 B2 -0.219699 0.01466680 52879 -14.97936 0  
 B3 0.712197 0.02243112 52879 31.75043 0  
 B4 1.649972 0.01185232 52879 139.21079 0  
 B5 0.490545 0.01054515 52879 46.51850 0

Correlation:  
 B0      B1      B2      B3      B4  
 B1 -0.182  
 B2 -0.433 0.501  
 B3 -0.381 0.015 0.000  
 B4 -0.544 0.281 0.572 -0.023  
 B5 -0.395 0.473 0.991 -0.002 0.518

Standardized Within-Group Residuals:  
 Min                 Q1                 Med                 Q3                 Max  
 -37.2585613 -0.4929532 -0.1104700 0.3622128 30.8658405

Number of Observations: 52889  
 Number of Groups: 5

\$SIINT  
                   B3  
 50.FALSE 0.054275483  
 80.FALSE 0.006830164  
 100.FALSE -0.031103205  
 50.TRUE 0.025956273  
 100.TRUE -0.055958716

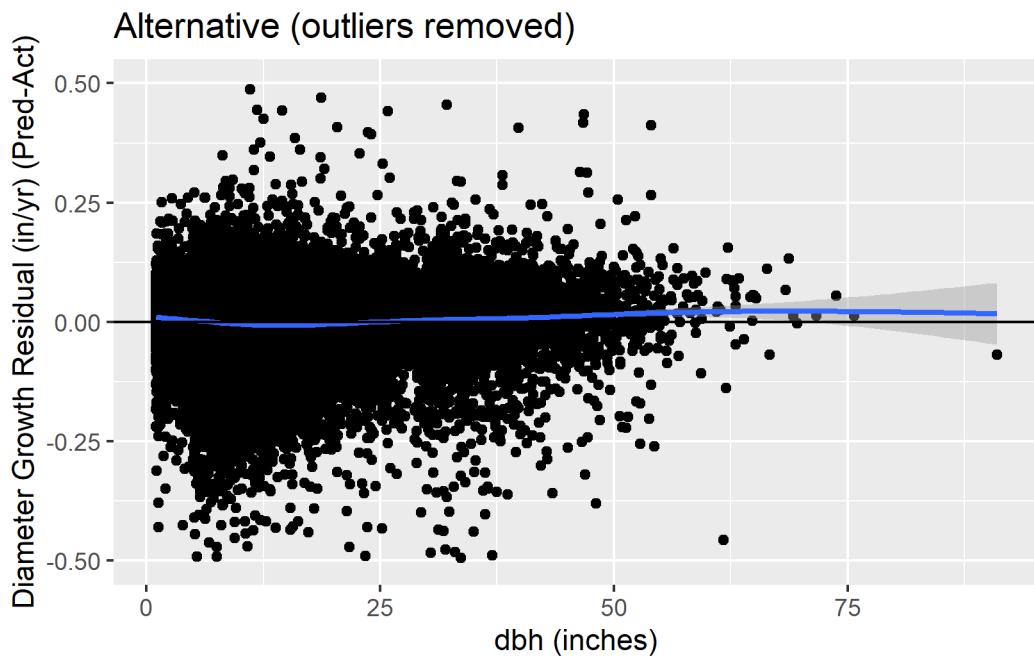
Residual Standard Error: 0.742584945600842 on 52879 degrees of freedom, AIC: 118654.5

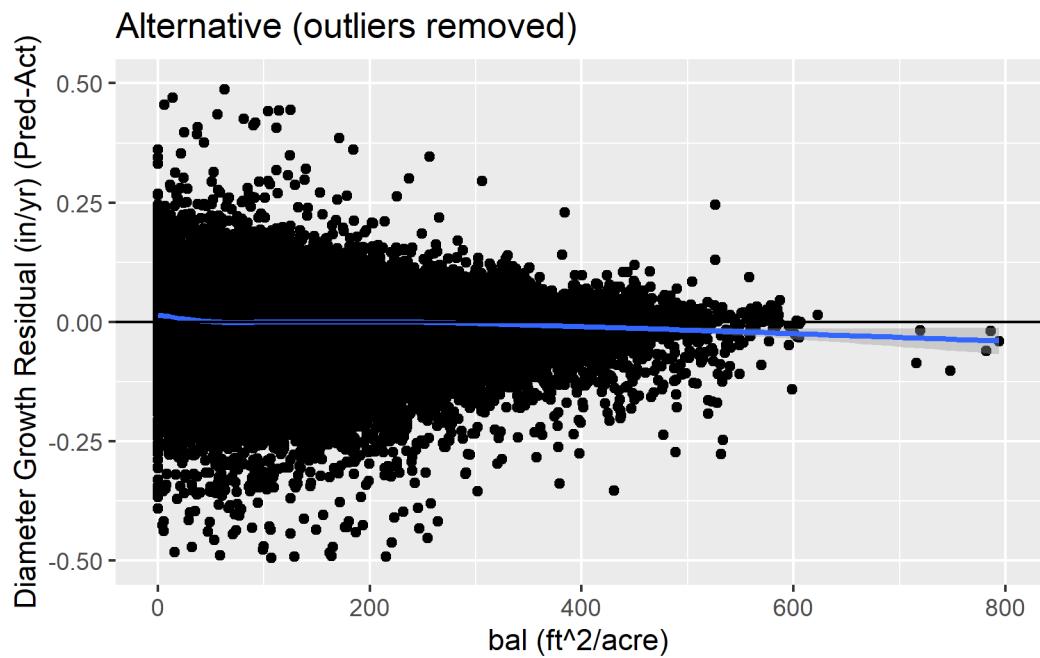
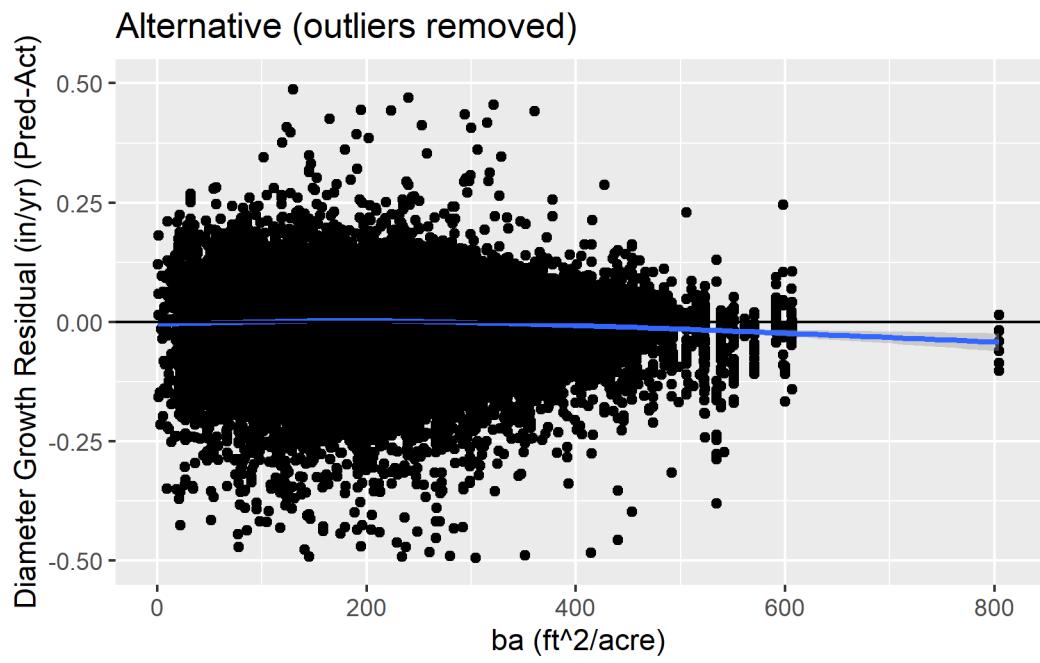
and for Equation 2:

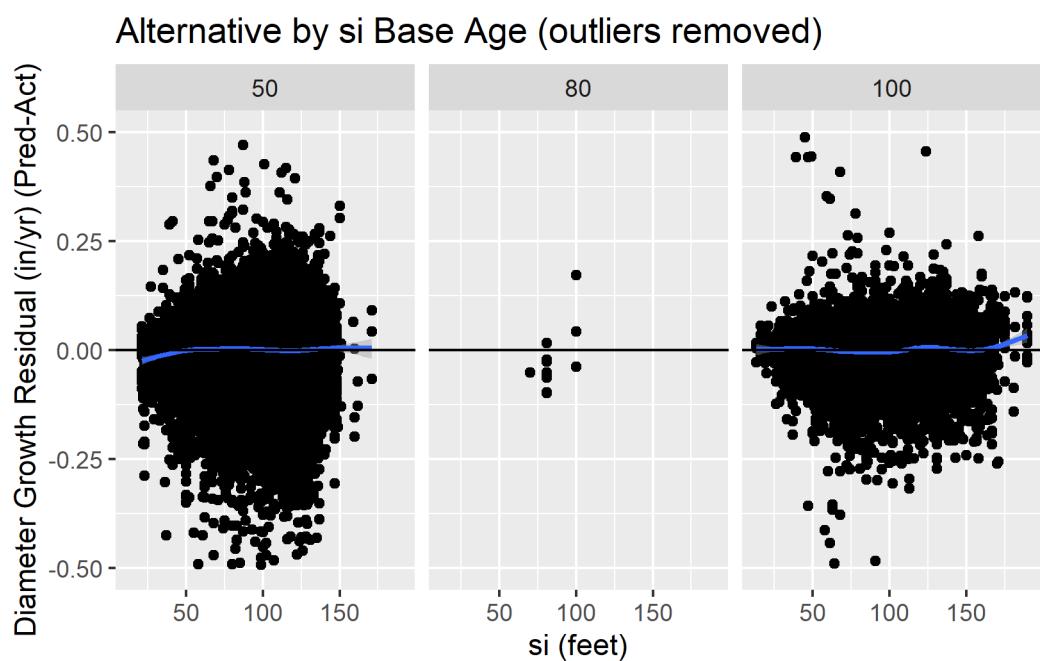
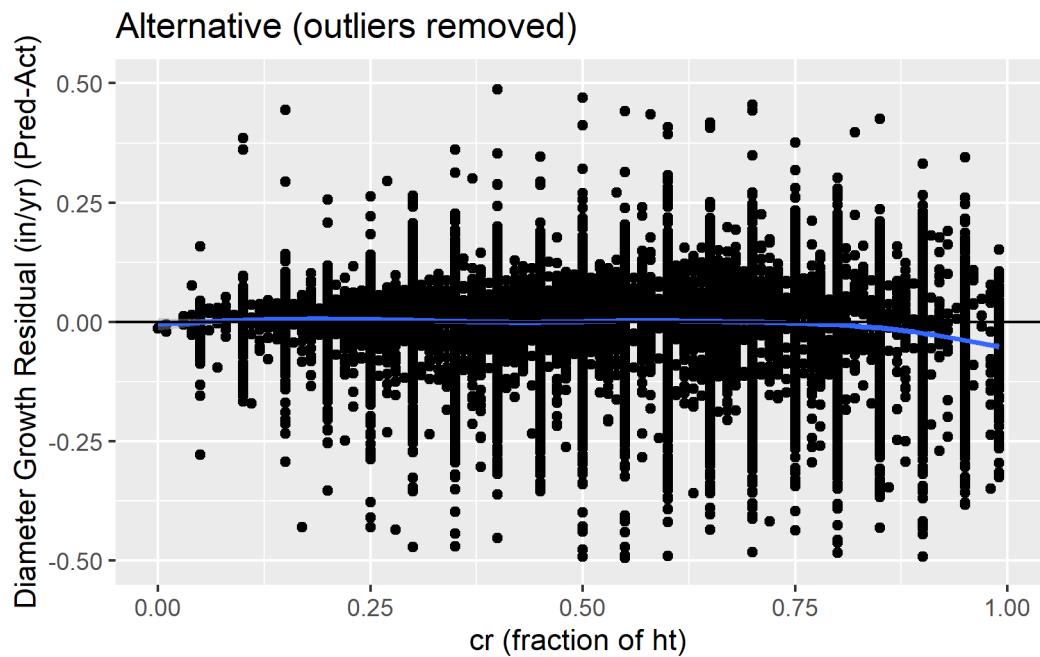
	Coef.	Std. error	t-stat.	p
B0	-2.1141719	0.0413919	-51.07694	0
B1	-0.7518152	0.00666531	-113.00300	0
B2	-0.2423855	0.0176098	-13.76421	0
B4	1.7549781	0.0115779	151.58035	0
B5	0.4716689	0.0113871	41.42126	0

Residual Standard Error: 0.796362713056519 on 52884 degrees of freedom, AIC: 126013.8

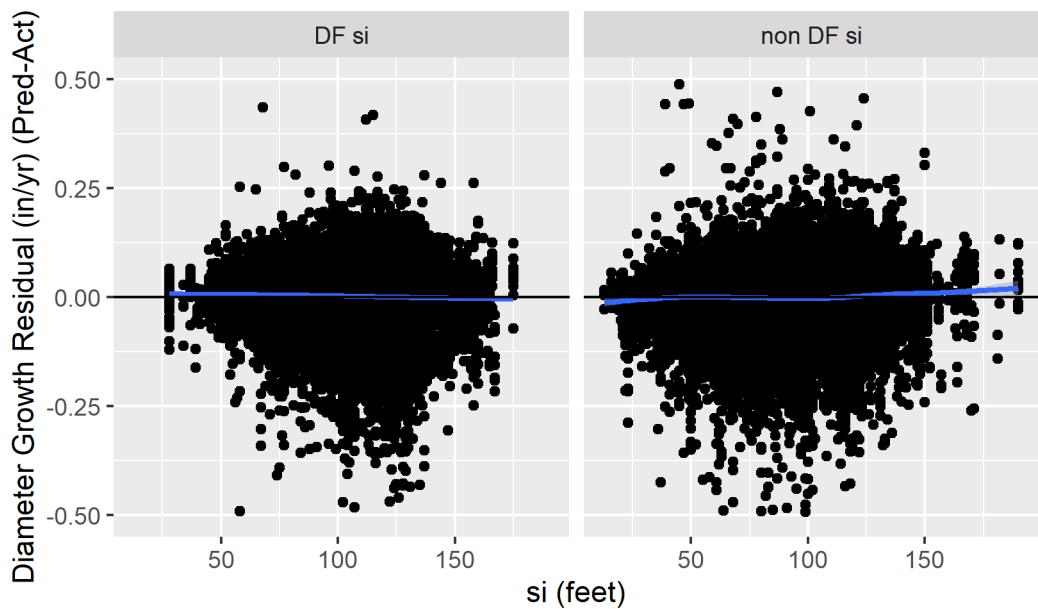
### Residual Analysis for Equation 1



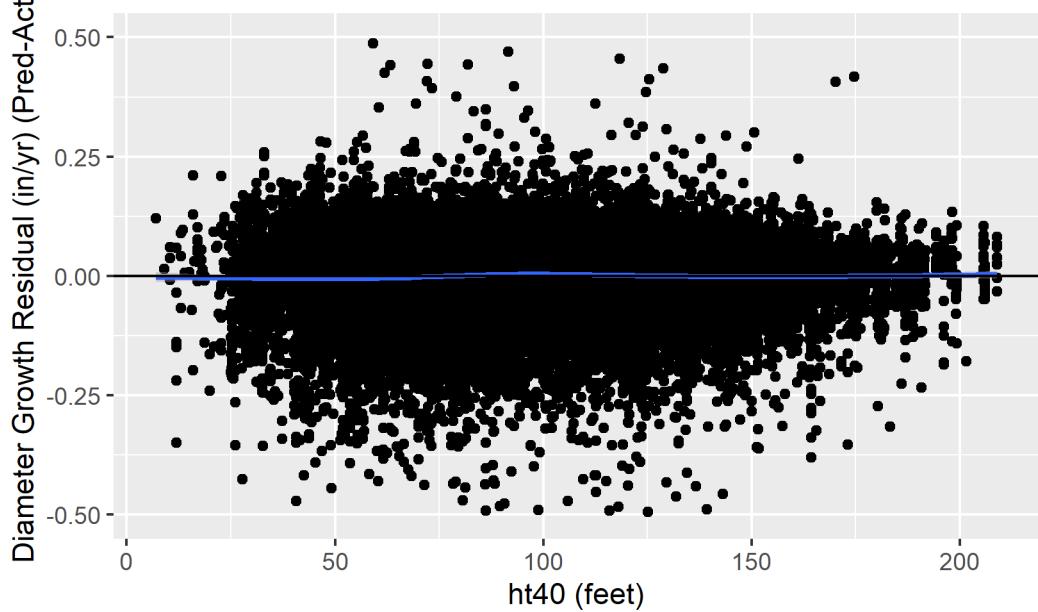


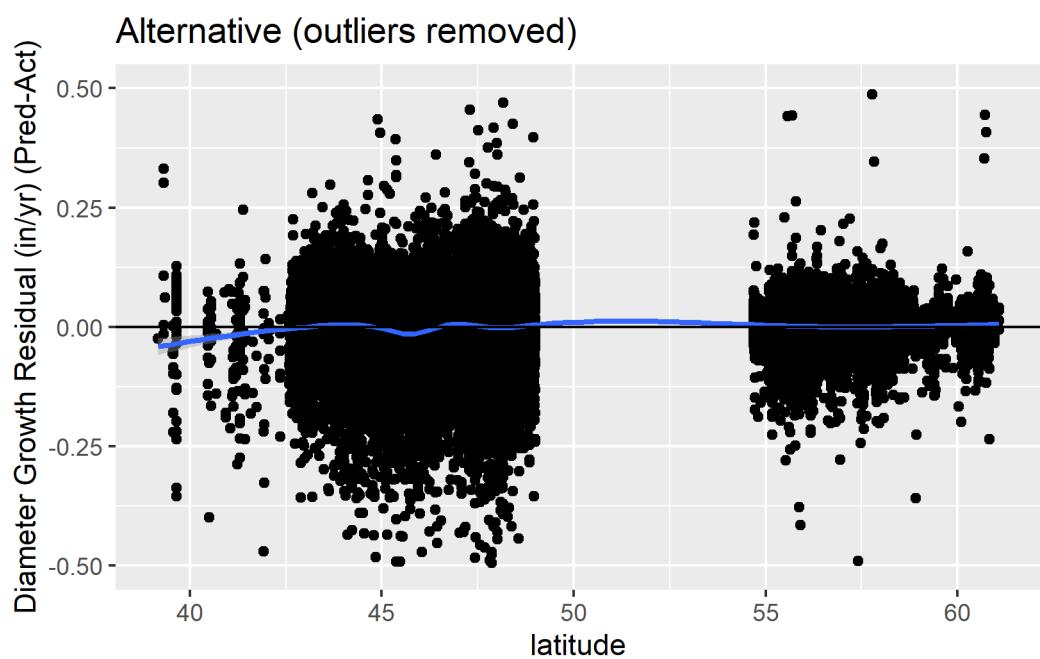
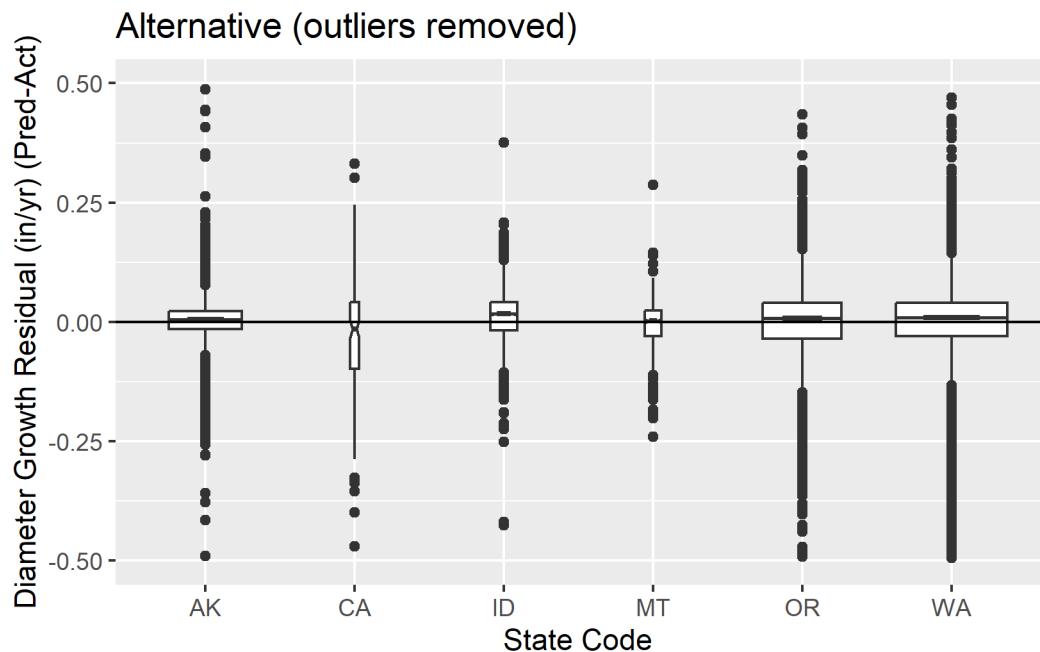


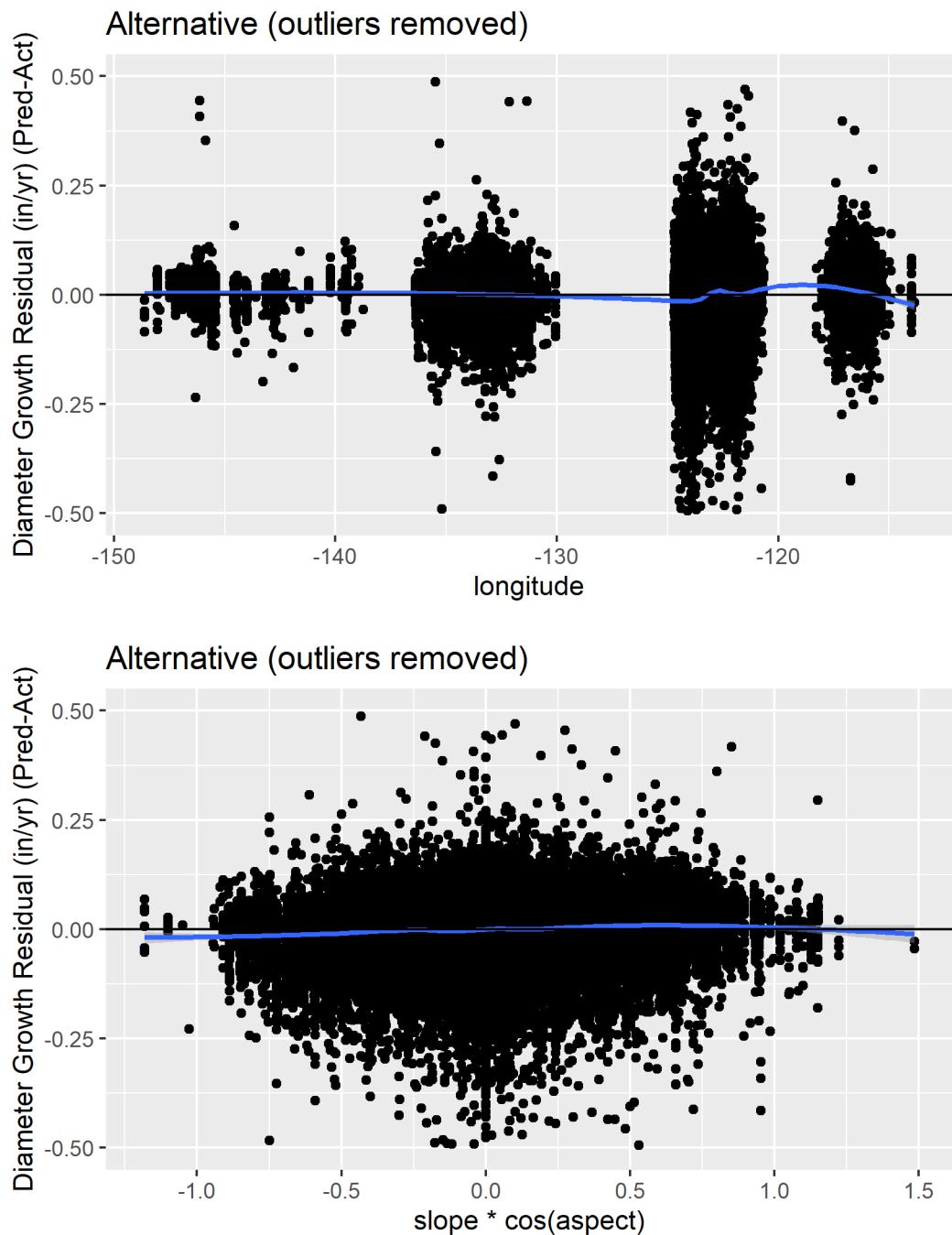
Alternative by si Species (outliers removed)

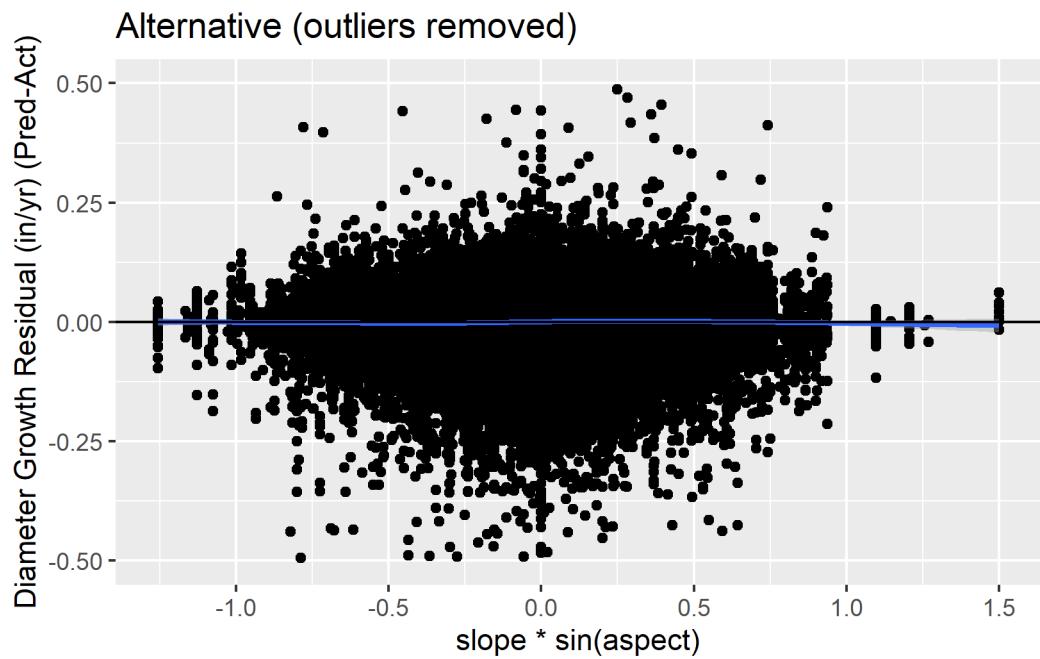


Alternative (outliers removed)

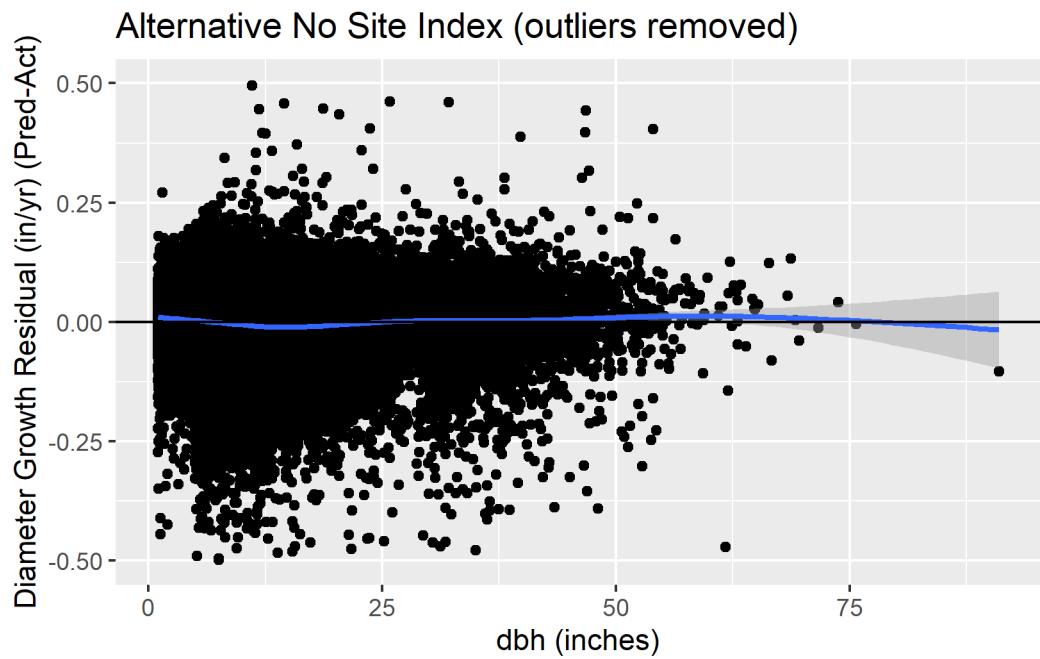


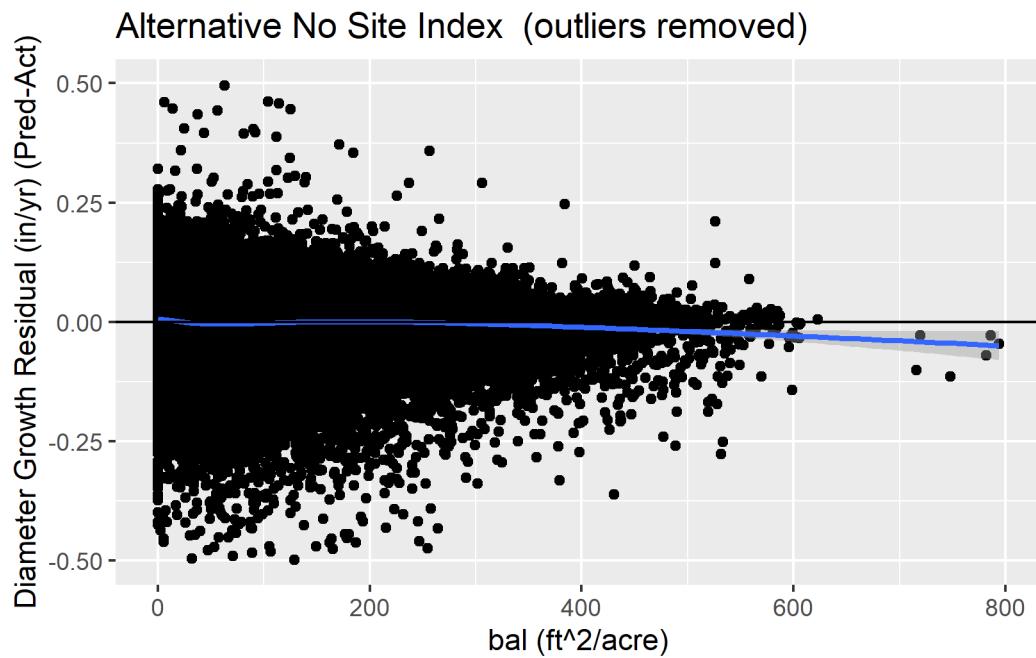
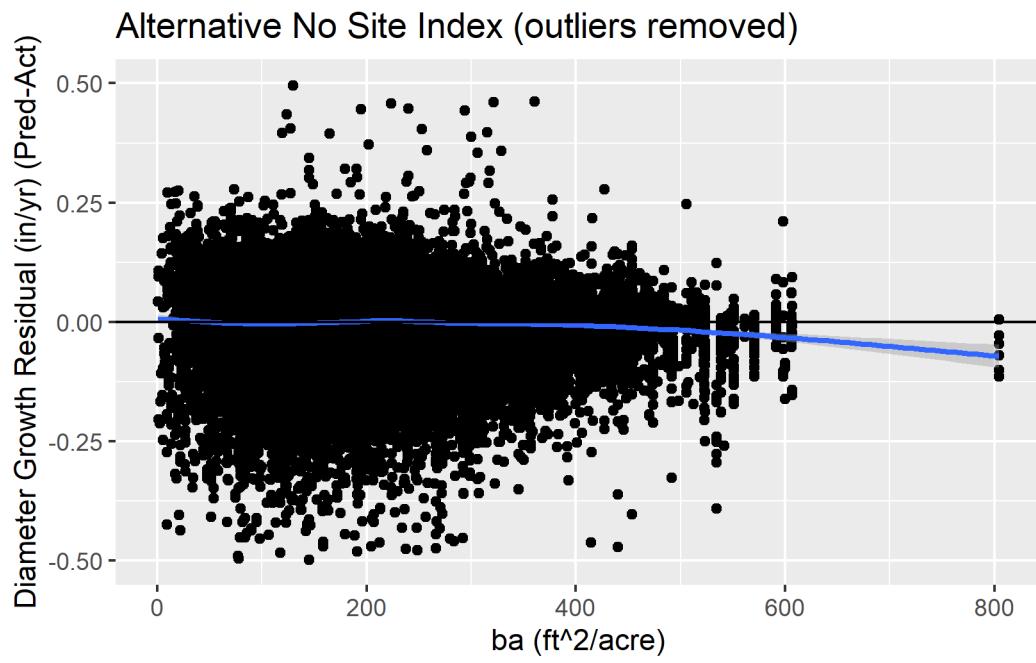


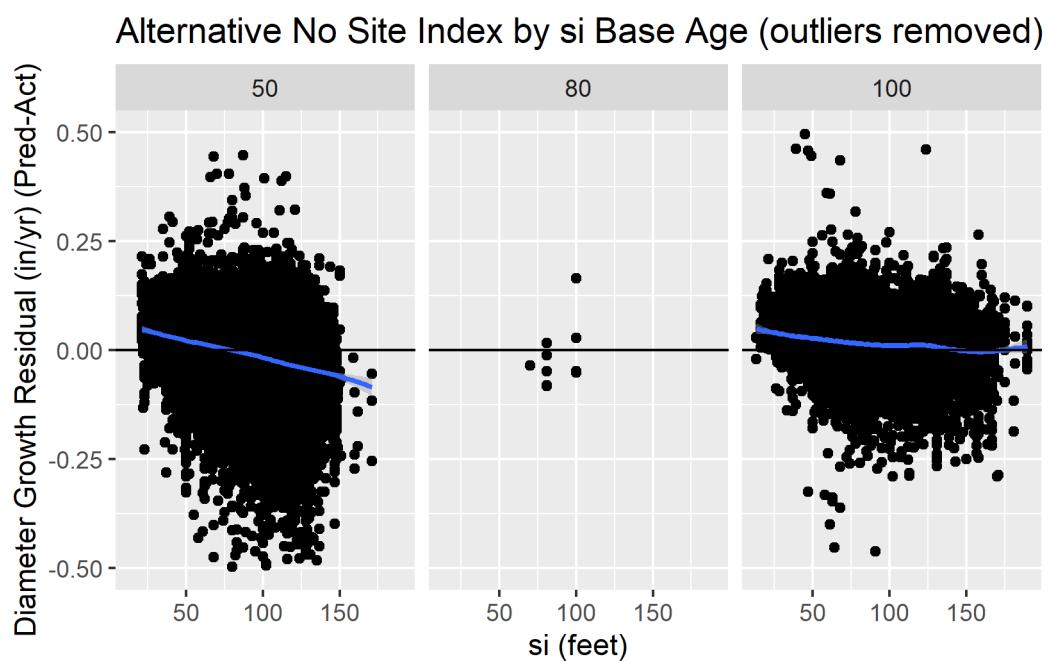
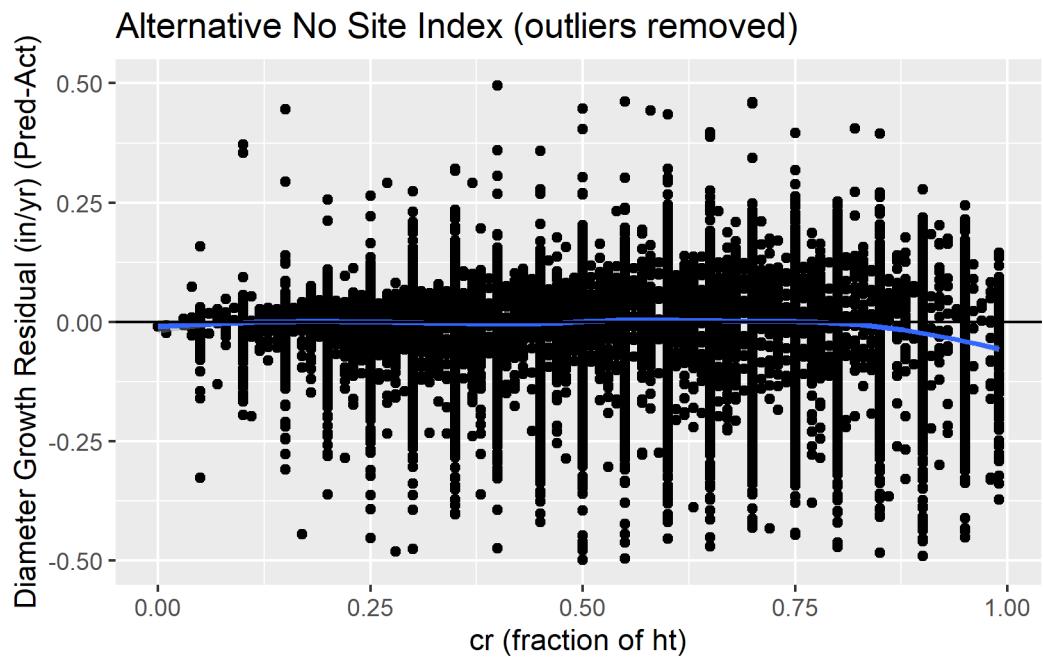




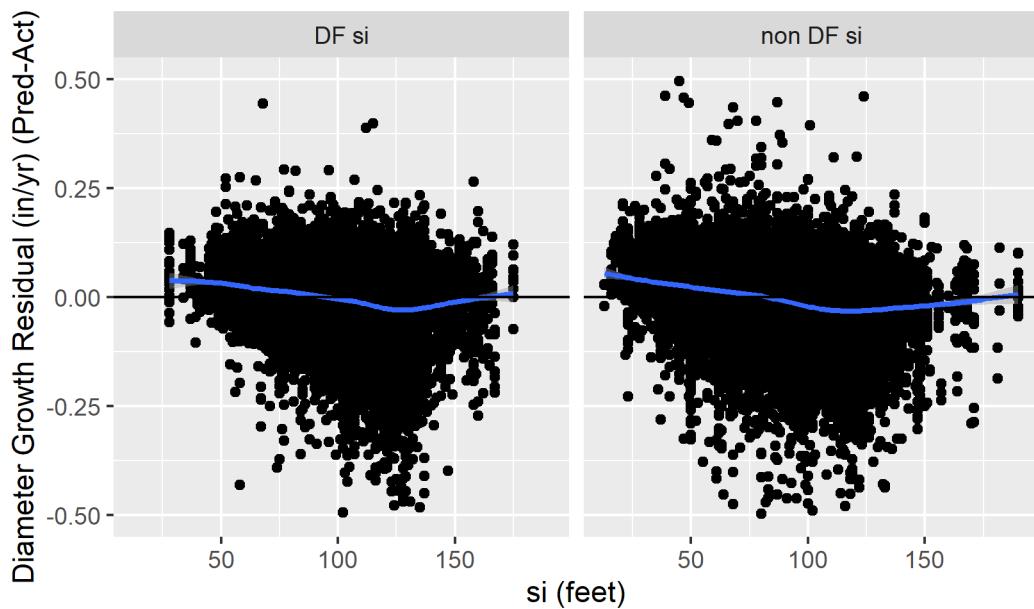
#### Residual Analysis for Equation 2



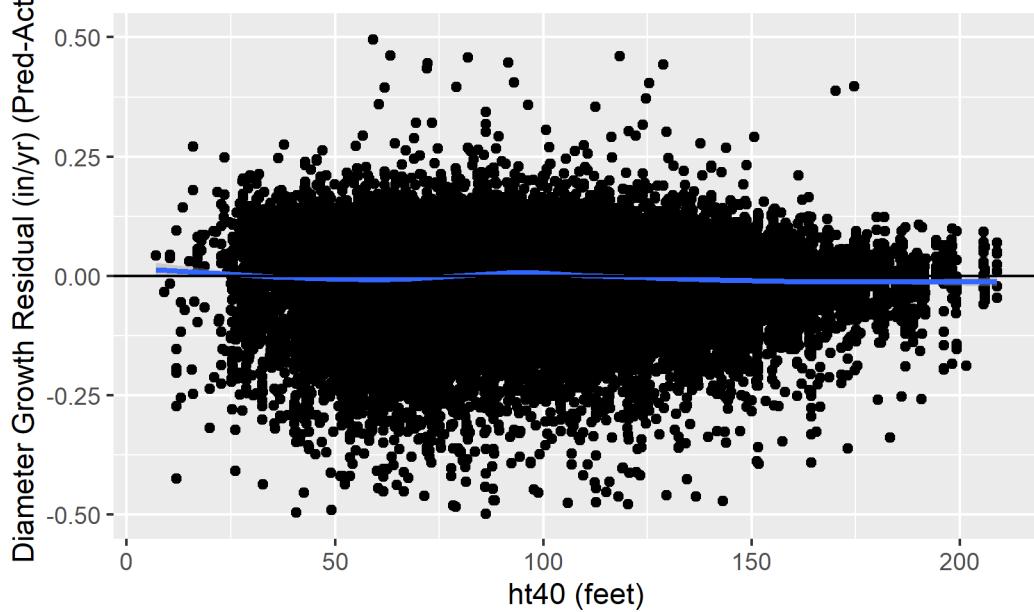


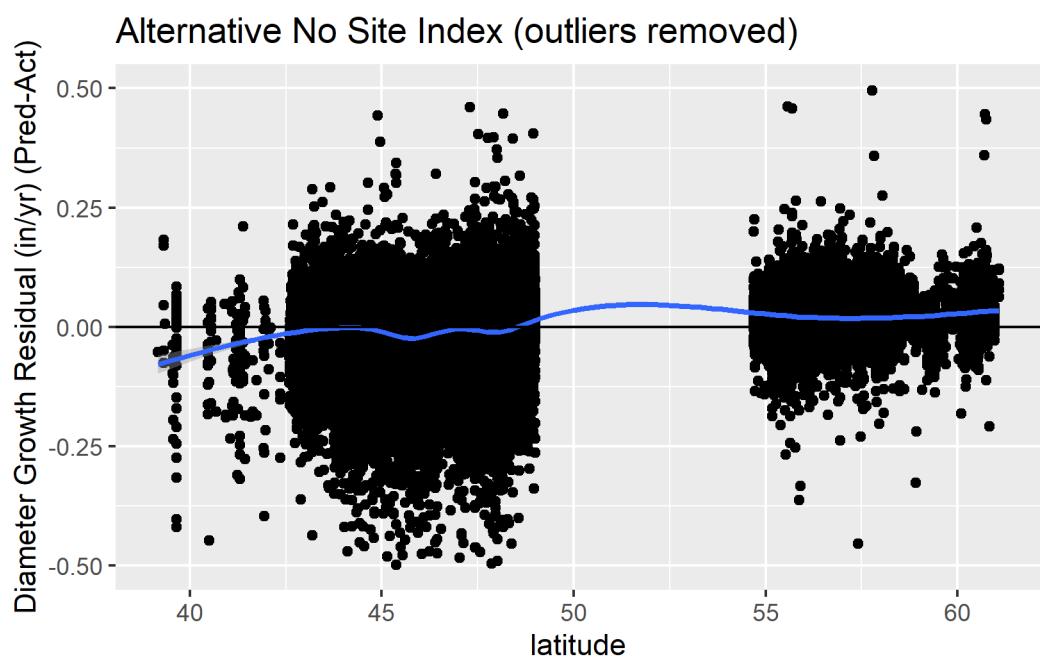
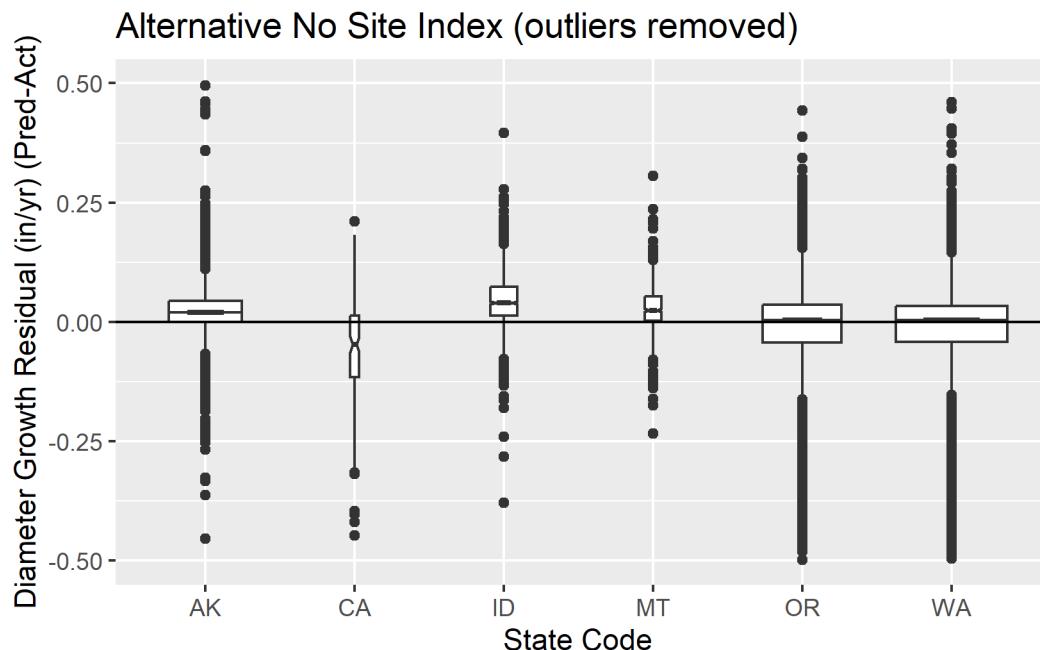


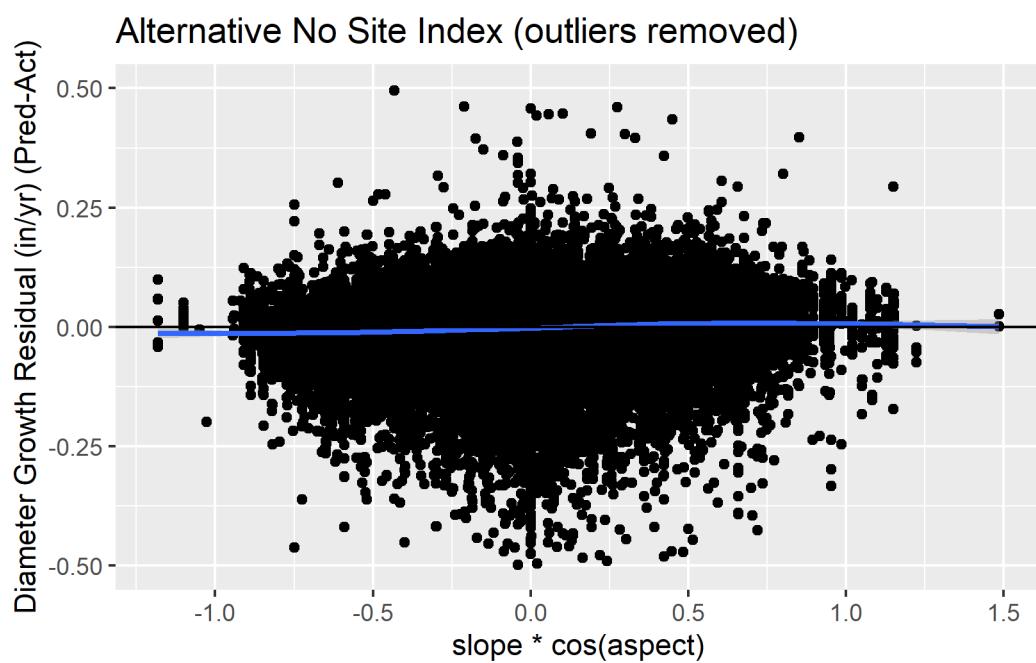
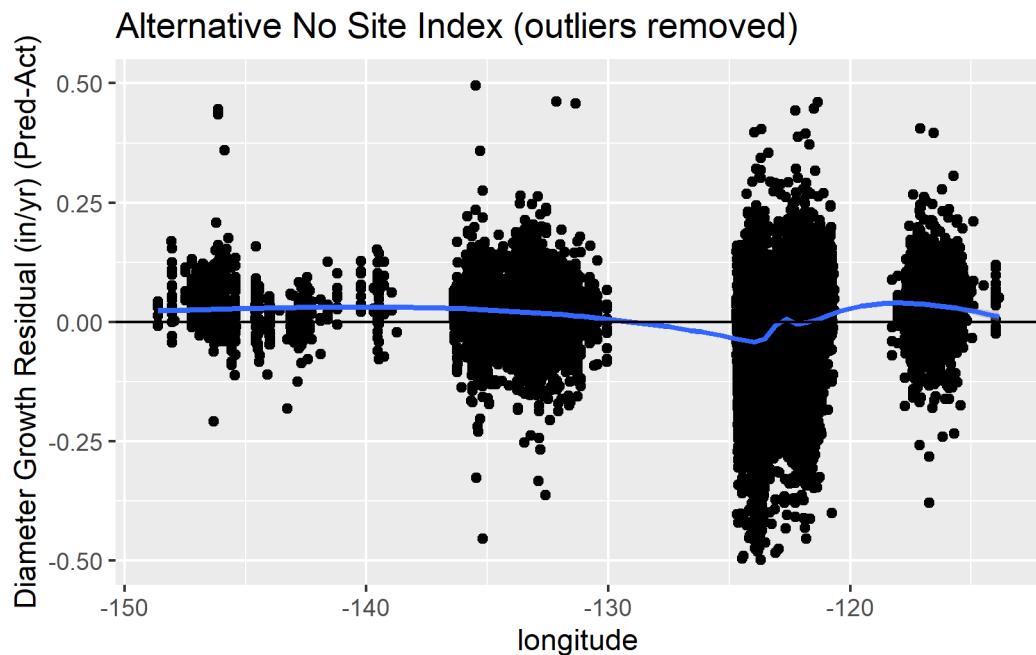
Alternative No Site Index by si Species (outliers removed)



Alternative No Site Index (outliers removed)







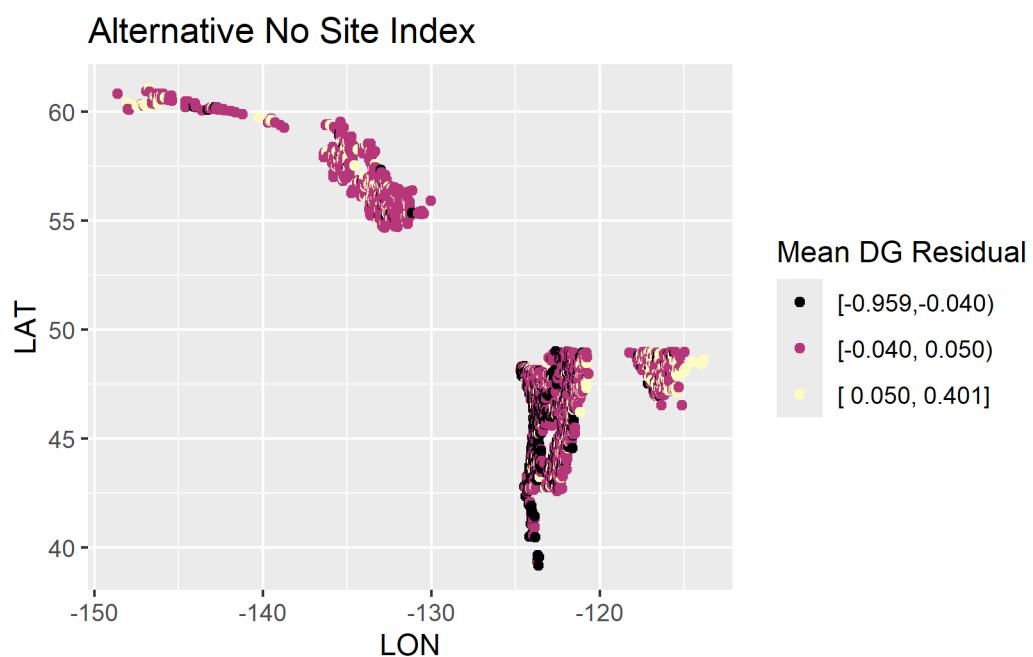
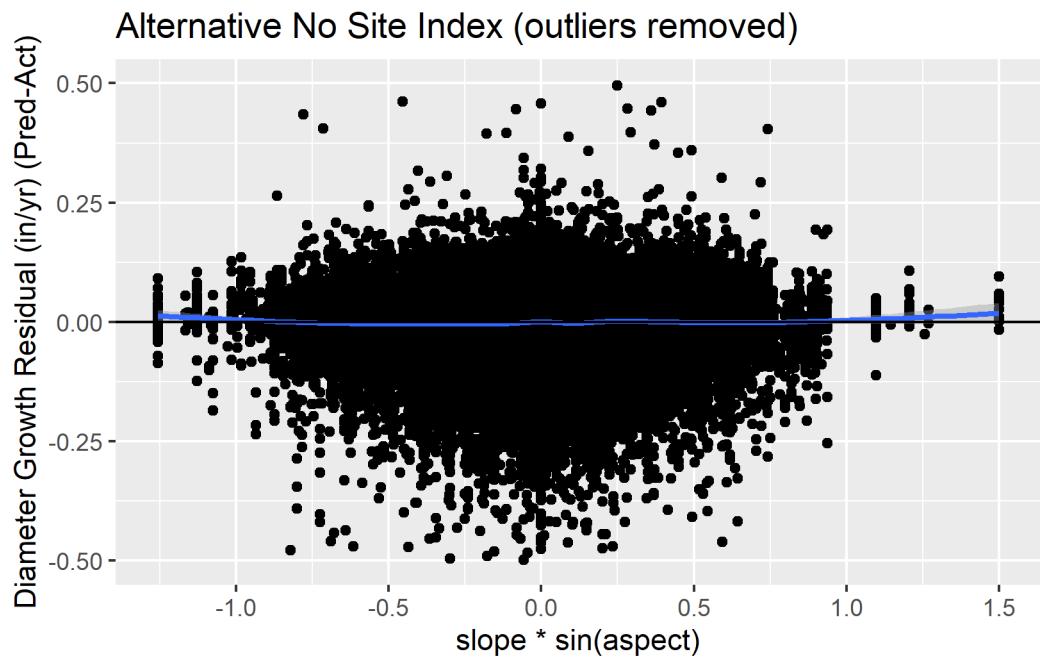


Table 3: Independent Variables for One Inch dbh Trees

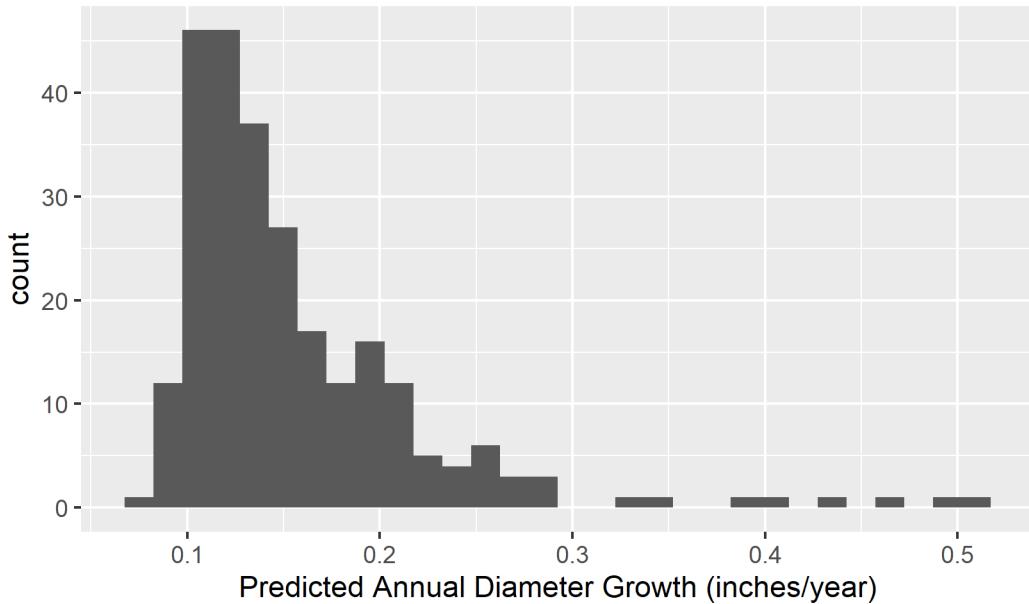
Variable	N	Mean	Std. Dev.	Min	Pctl. 25	Pctl. 75	Max
ba	255	180	92	5.4	109	241	523
bal	255	179	92	5	108	240	523
ht	255	9.5	2	5	8	10	20
cr	255	0.45	0.26	0.05	0.25	0.64	0.99
si	255	85	31	18	60	107	171

## Discussion

Removing `si` degrades the fit significantly. There is an over-prediction bias for the eastern states for both equations, but is mitigated by the `si` term in Equation 1. The residual trend with longitude confirms this.

### Equation Behavior for Very Small Trees

#### Equation 1 Predictions for Trees with One Inch dbh Trees



## Equation 2 Predictions for Trees with One Inch dbh Trees

