

Formulating climate sensitive yield projections using transfer functions

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Yield Projections

- Silvicultural planning
- Supports Allowable Annual Cut (AAC) determinations
- Carbon Budget Monitoring
- National wood supply
- ...many other applications

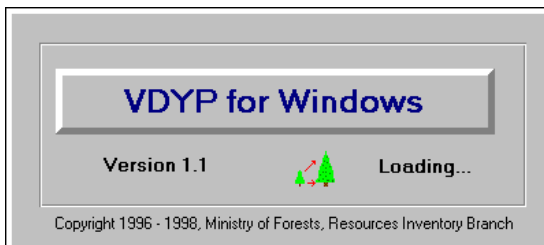
Yield Projection Systems and Climate

- **Why the need for climate variables within G&Y models?**
 - Climate change is accelerating
 - Longer projection periods
 - Climate affects productivity (30% of variability in Site Index due to climate^a)
 - CBST
 - Blue Ribbon Panel Report (2018); Penner Report (2021); Fletcher Report (2023)
- **BC models: climate static**
 - AB*, SK, and MB models climate static (*climate vars added to MGM)
 - ON – climate sensitive site index curves
 - NB (NS, PEI) and QC – some climate sensitive functions

^a Monserud et al. 2006

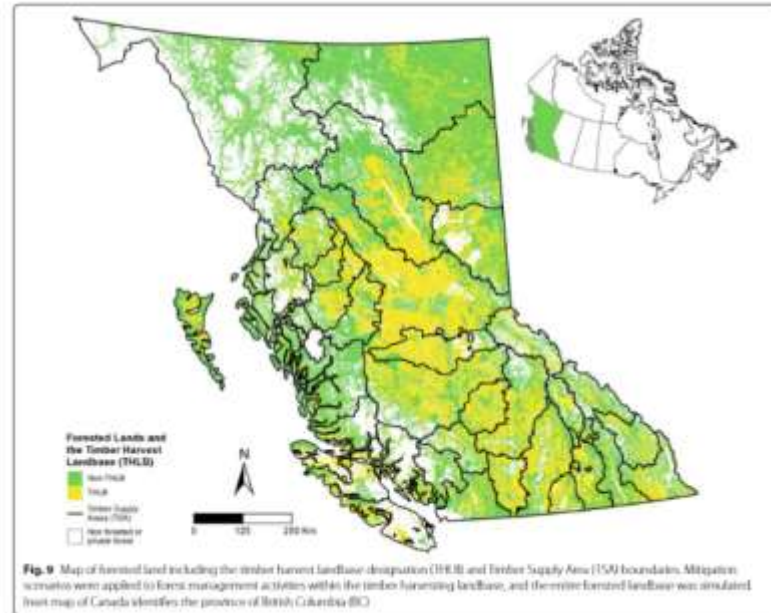
Yield Projection Systems in BC

Unmanaged stands



Stand-level
Crown closure, Basal area, Site Index

Pure and mixed-species
TSR and inventory updates
Deterministic



Smyth et al. 2020

Managed stands



Individual tree (TASS)
Mixed-species (TASS III)

Meta-model (TIPSY)
Stand-level (TIPSY)
TSR (TIPSY)



TASS

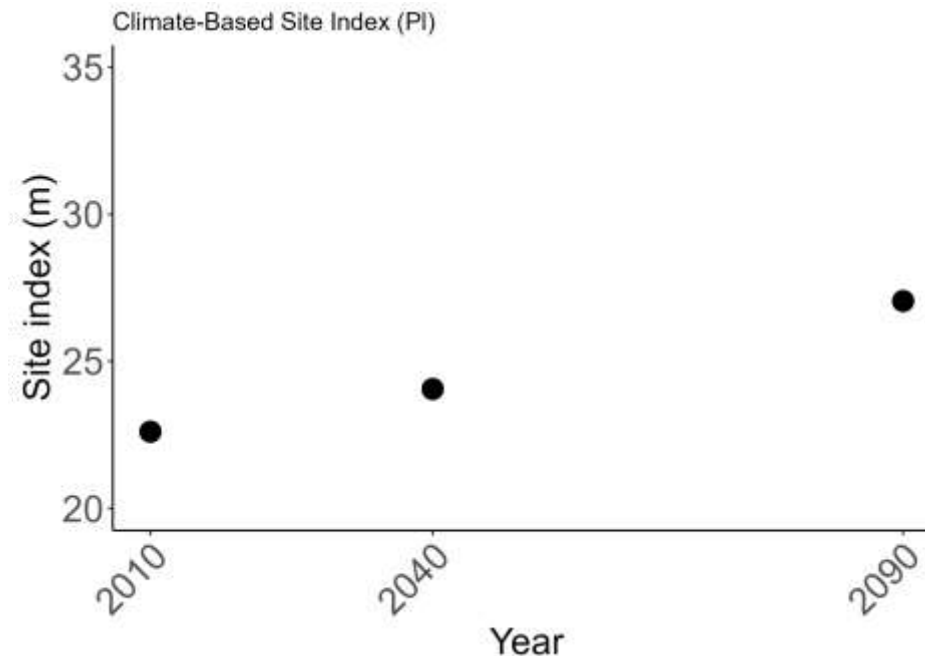
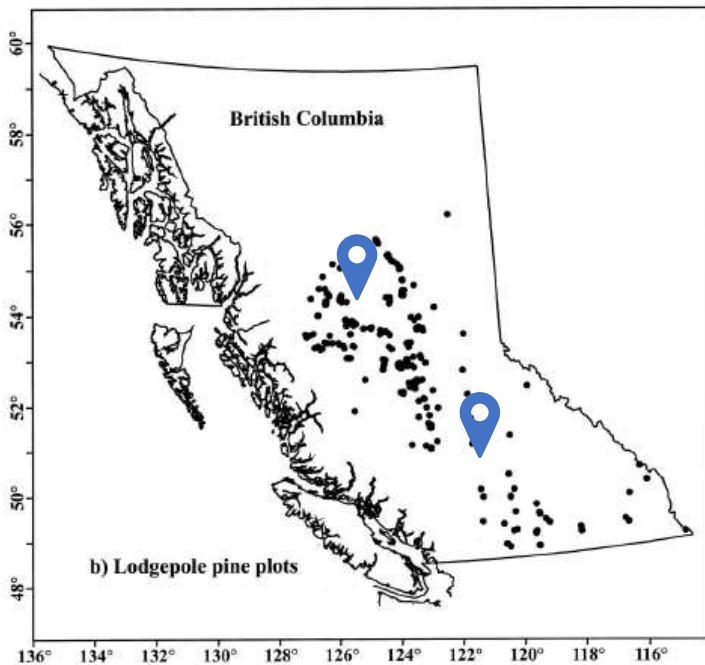


TIPSY

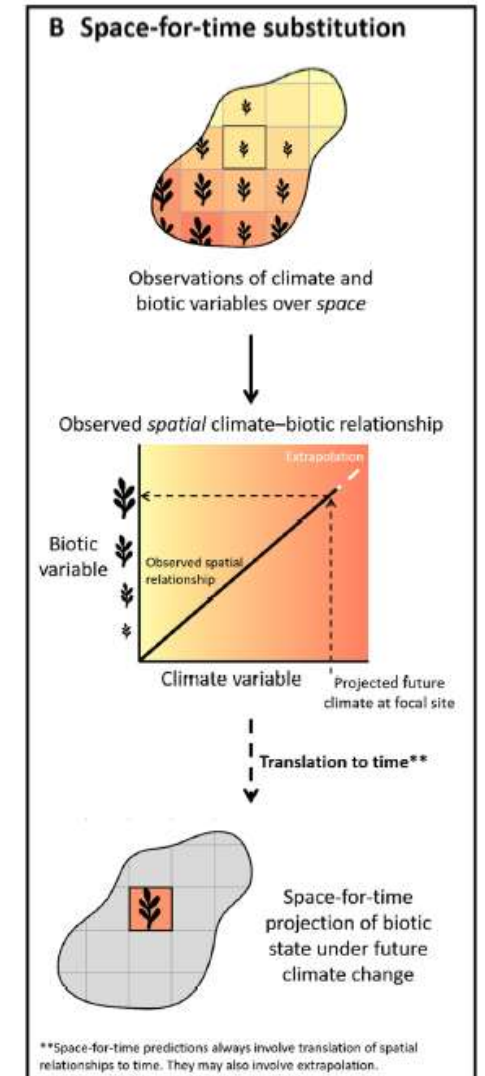
How to incorporate climate effects into G&Y?

1. Temporary Sample Plots (Space-for-time substitution)

- Sample population = locally adapted trees
- Most studies predict increased site productivity
- Nigh et al. (2004) for site index [SI = f(Climate Vars.)]



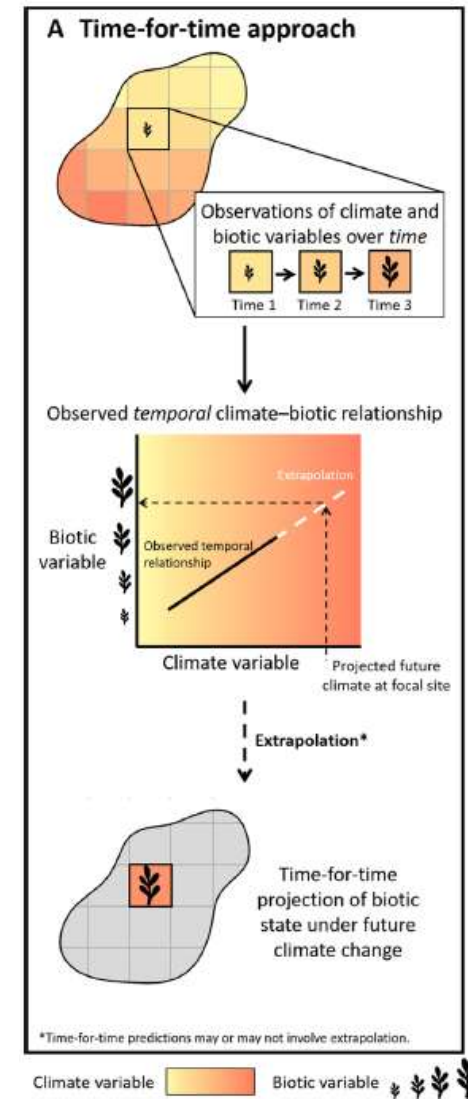
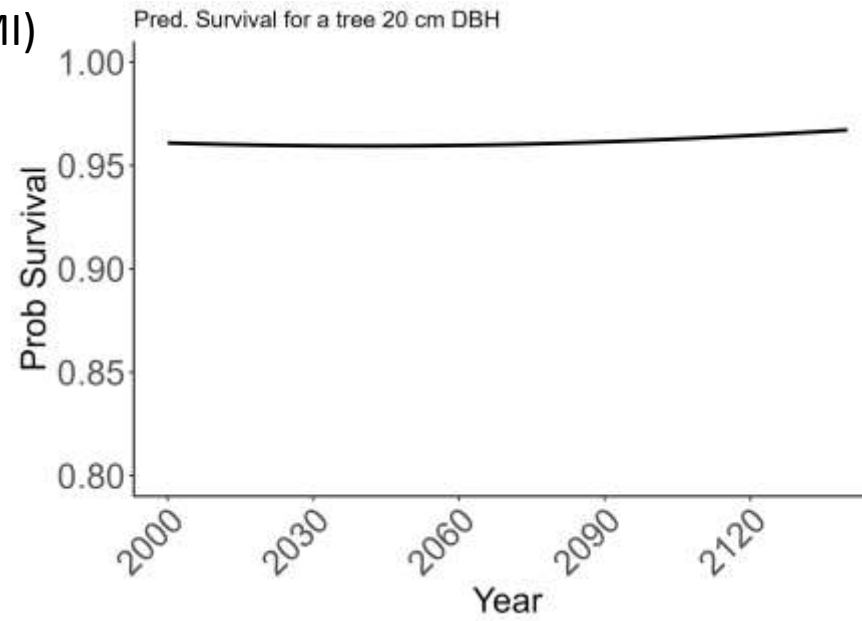
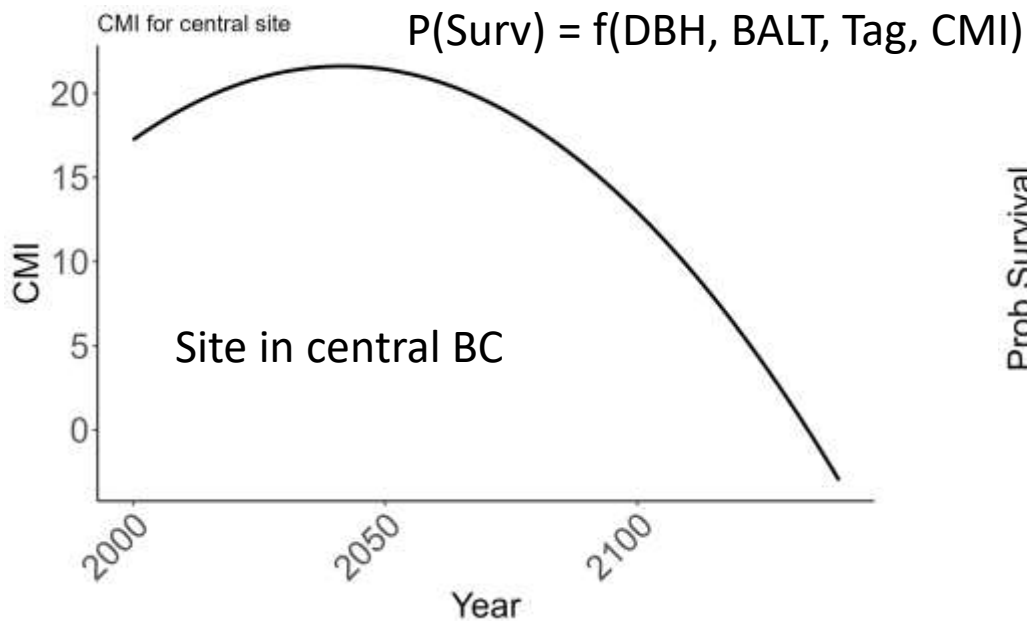
Model
from Nigh
(2004)



How to incorporate climate effects into G&Y?

2. Permanent Sample Plots (time-for-time)

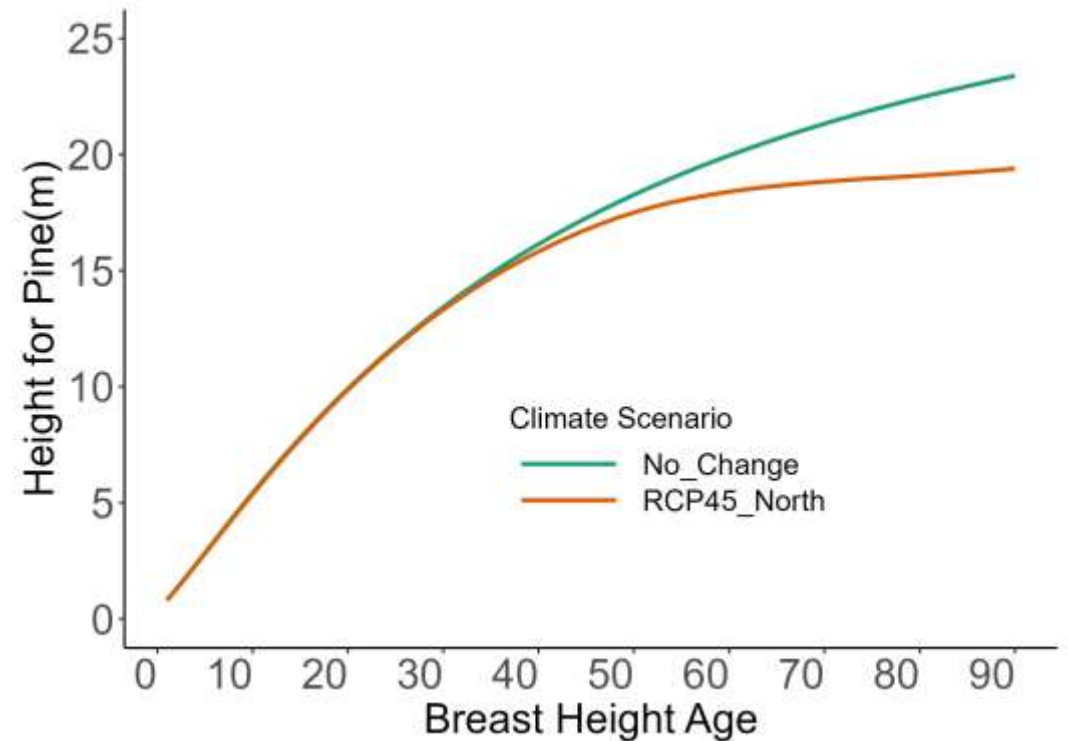
- Sample population = locally adapted stands
- “Marginal effect” of climate is usually small
- Oboite and Comeau (2020) for diameter growth, Cortini et al. (2017) for survival in MGM



How to incorporate climate effects into G&Y?

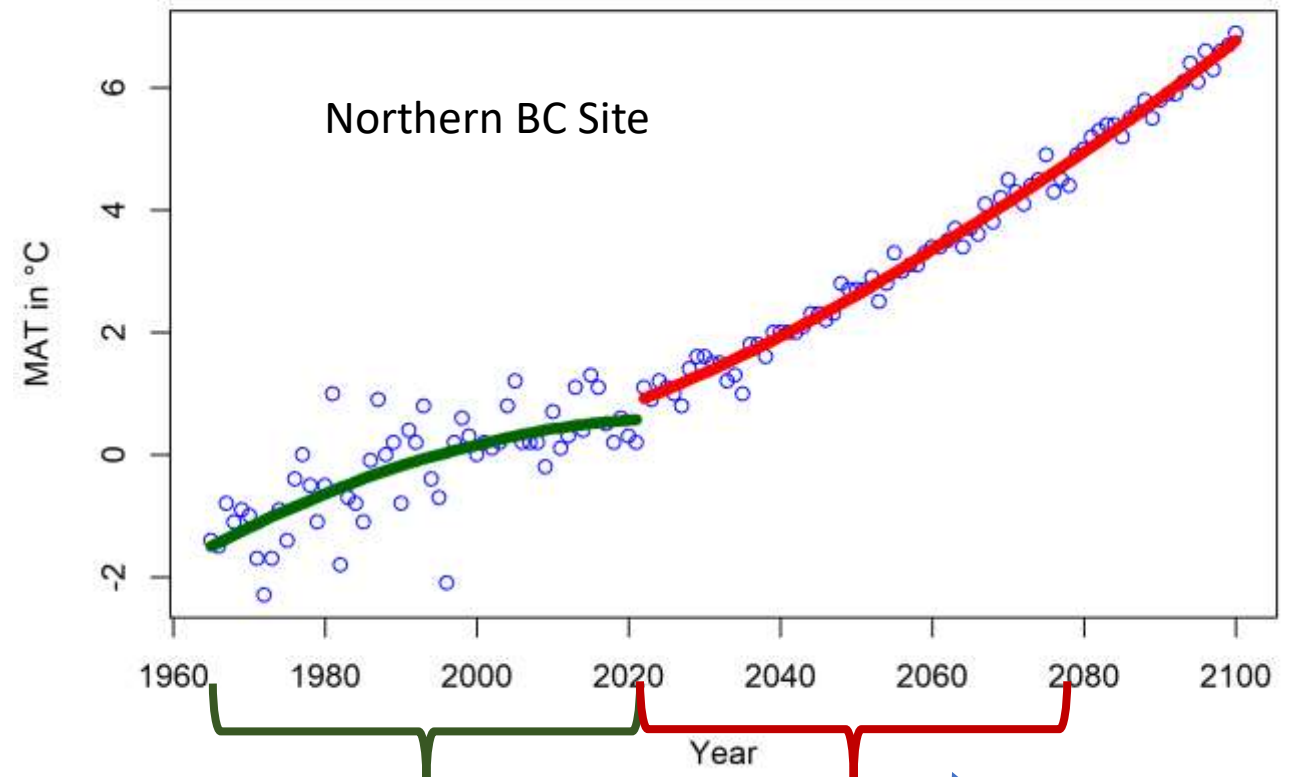
3. Stem analysis

- Sample population = locally adapted trees
- Uncertainty around sample tree history
- Sharma et al. (2015) for Height-age (Site index) curves



How to incorporate climate effects into G&Y?

- Approach 2 requires large extrapolation
- Future migration lag is underestimated
- Not designed to accommodate seed transfer
- Ignore genetic-by-environment interaction

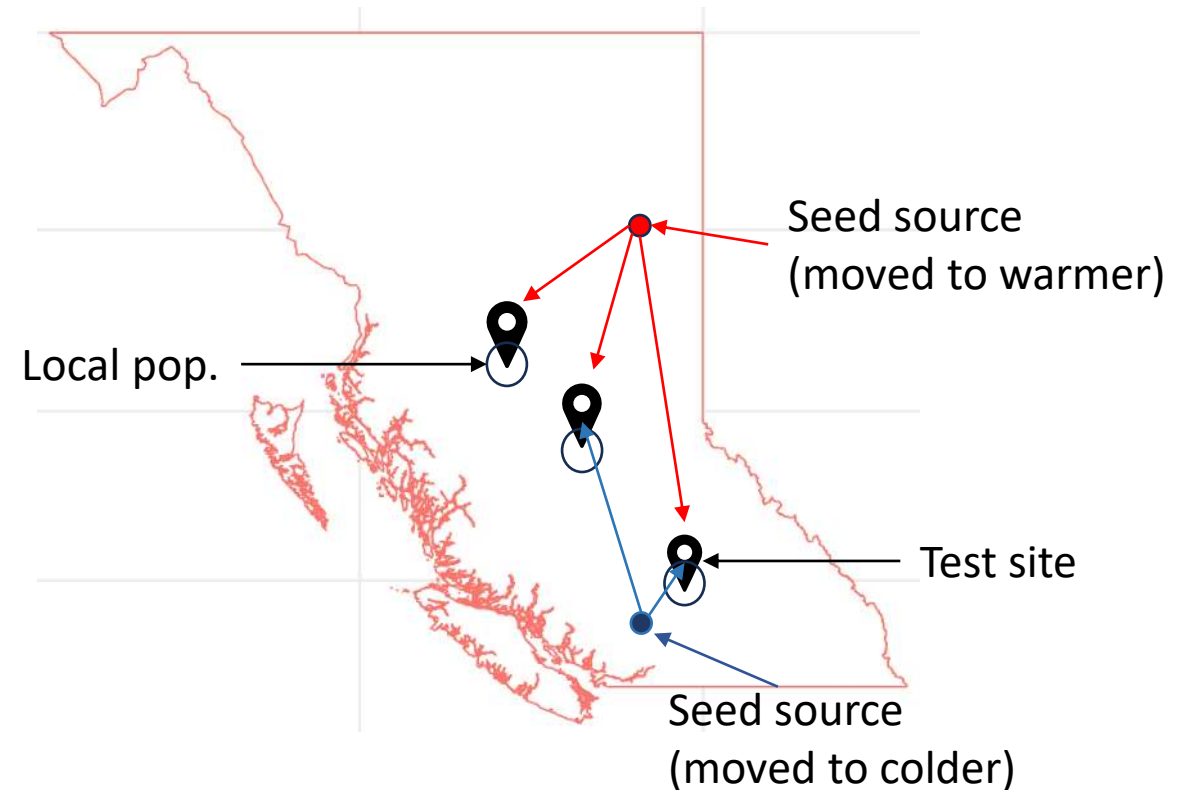


delta MAT ~ 1°C

delta MAT ~ 3.5°C

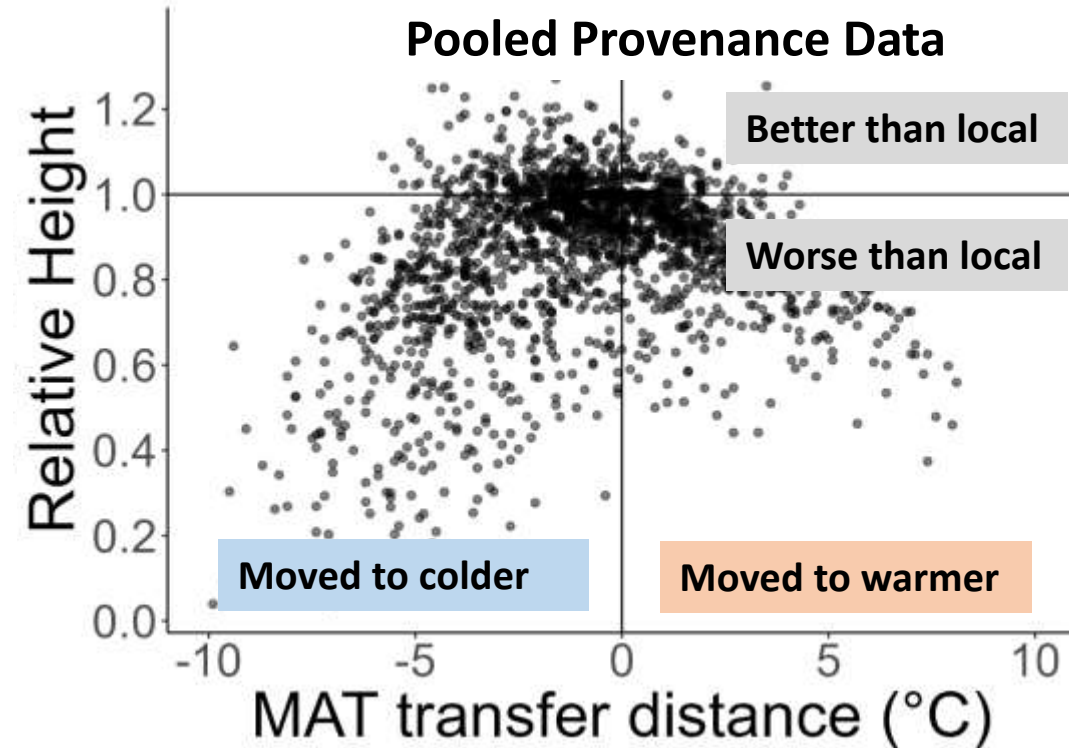
Transfer Functions and Climate Change

- TFs relate attributes (e.g., HT, Survival) to climate transfer distance



Transfer Functions and Climate Change

- TFs relate attributes (e.g., HT, Survival) to climate transfer distance
- Population-specific response to climate change
- Accounts for climate-maladaptation
- Less extrapolation
- **How to link to G&Y models?**
- **Of limited use within TSR**



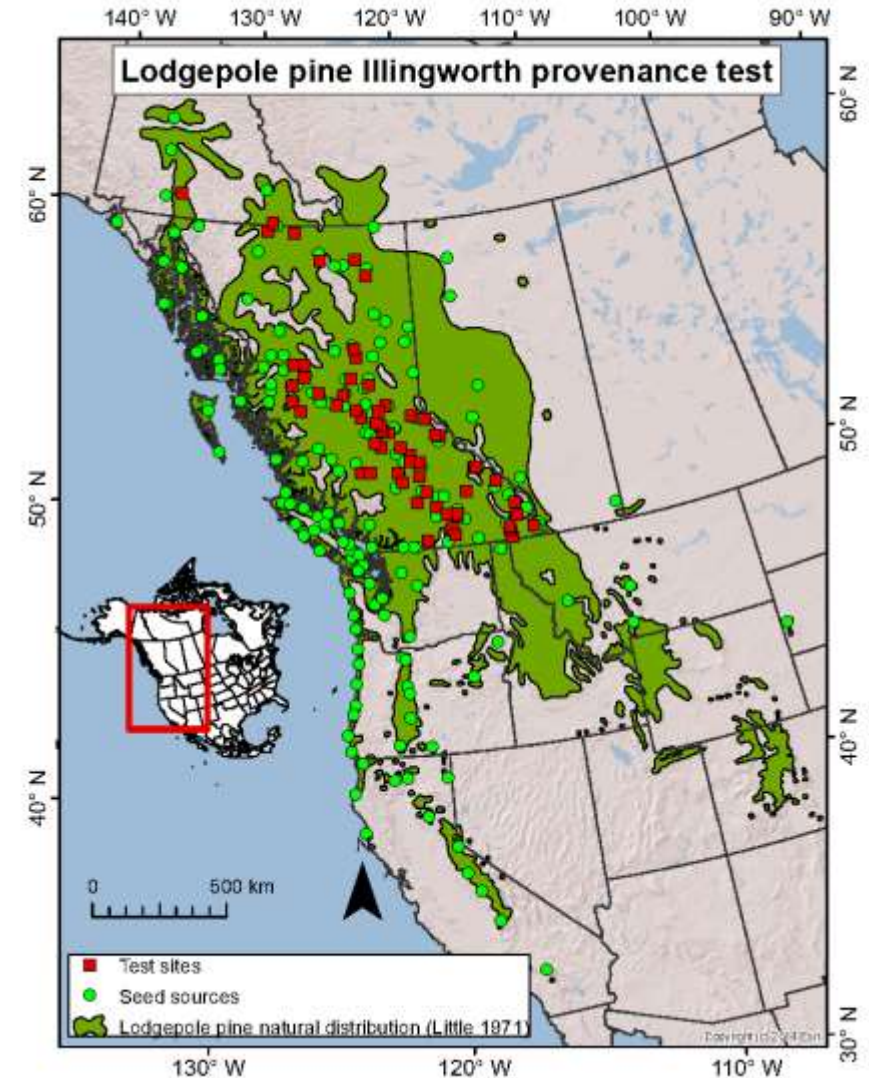
Transfer functions in Growth and Yield Modeling

O'Neill and Nigh (2011): Adjust Height-Age curves for Lodgepole Pine using Transfer Function for Relative Height

- Illingworth Data (est. 1974, 60 sites, 142 pops.)
- Tested in TASS

Objectives:

1. Revisit TF for Height
2. Formulate a TF for Survival
3. Link both the TF for Height and TF for survival to a GY model

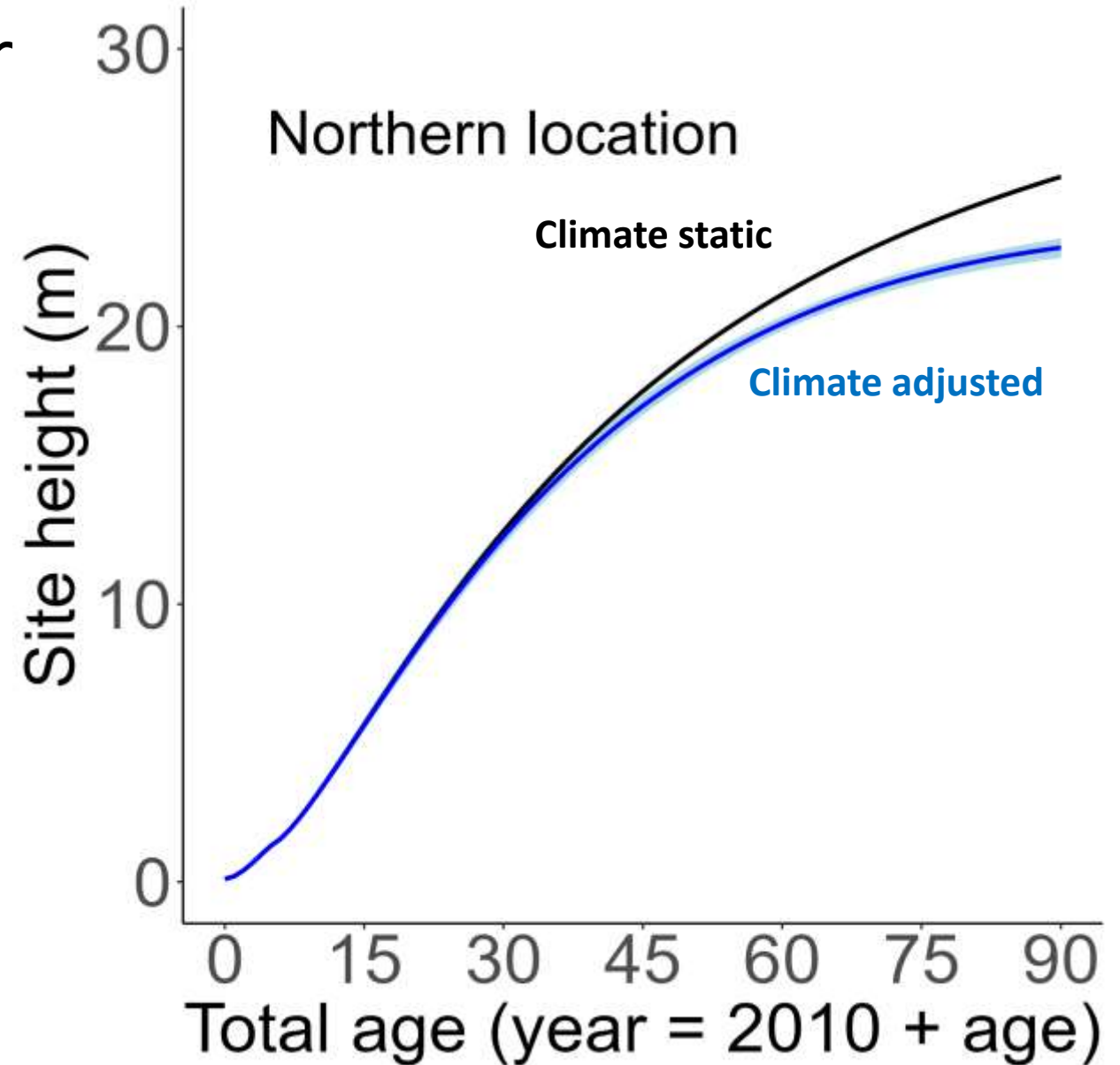


Source: Amy Vallarino

Using TFs to modify gr

- **Revised TF for Relative Height (Hp)**

- Includes a population-specific interaction term
- **GLM** ($Y \sim N(u, \sigma^2)$), where:
($g(\mu) = \ln(\mu)$)
- Relative Height (Hp) used as a multiplier in Height-Age equations



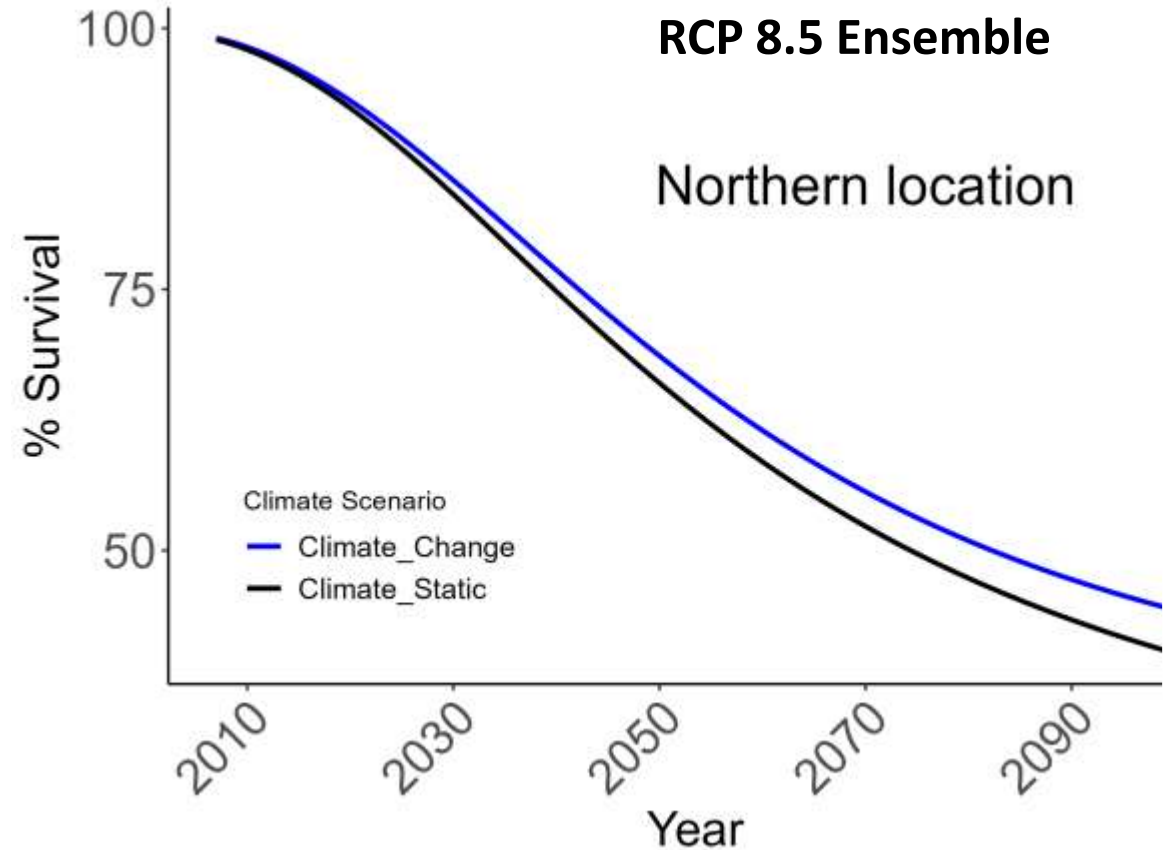
Using TFs to modify growth models

- **TF for Survival**

- Dynamic over time
- Model fit using repeated measures from Illingworth Trial
- Genetic-by-environment interaction

- **Tree Survival in TASS**

- Mortality controlled by neighbourhood, crown size, random events
- Too complex



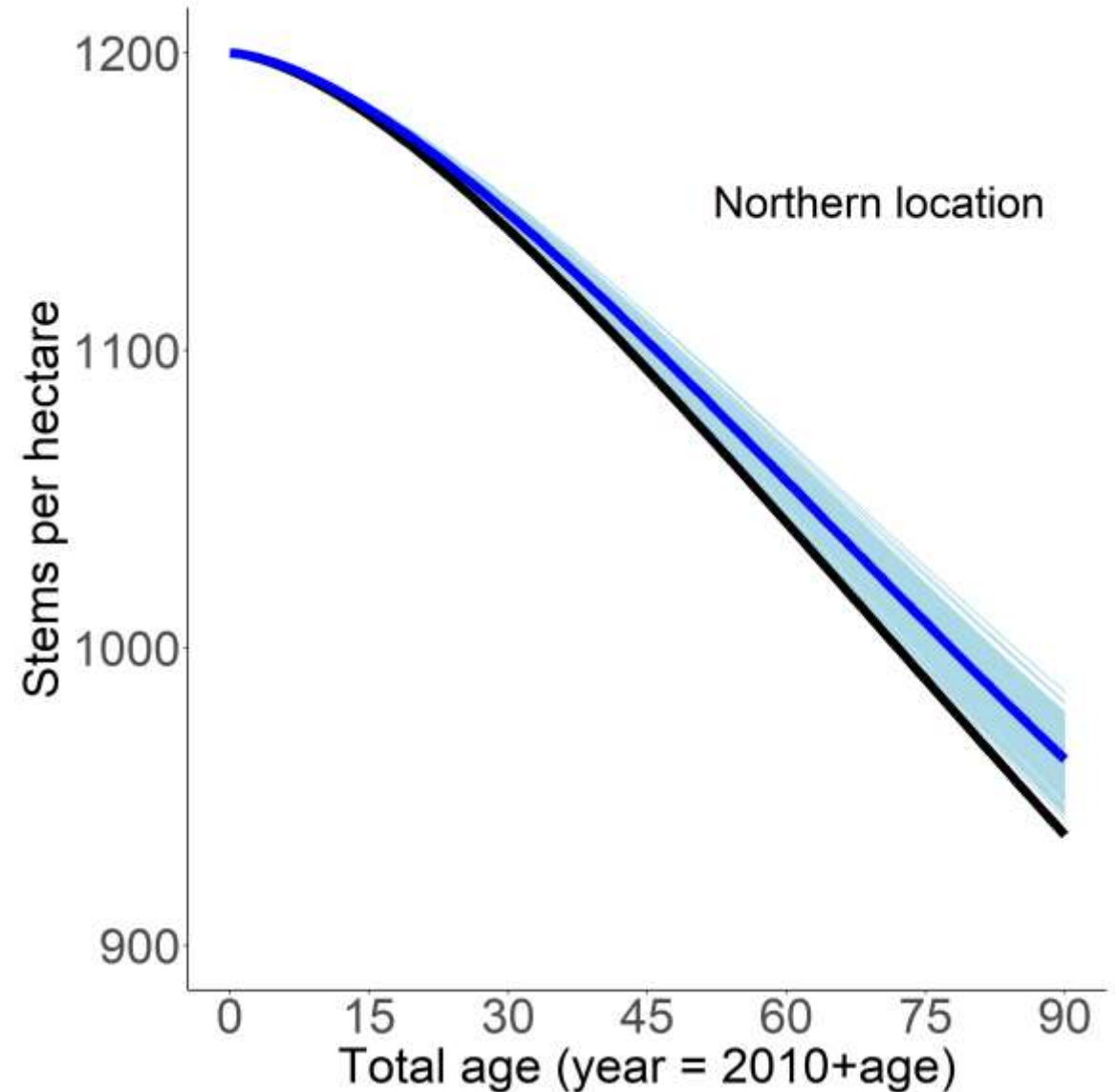
Using TFs to modify growth models

- **Meta-model of TIPSy:**

- Volume (m^3/ha) = $f(\text{Site Height}, N)$
- Site Height = $f(t, H_0, t_0)$
- $N_2 = f(t_2, t_1, N_1, SI)$

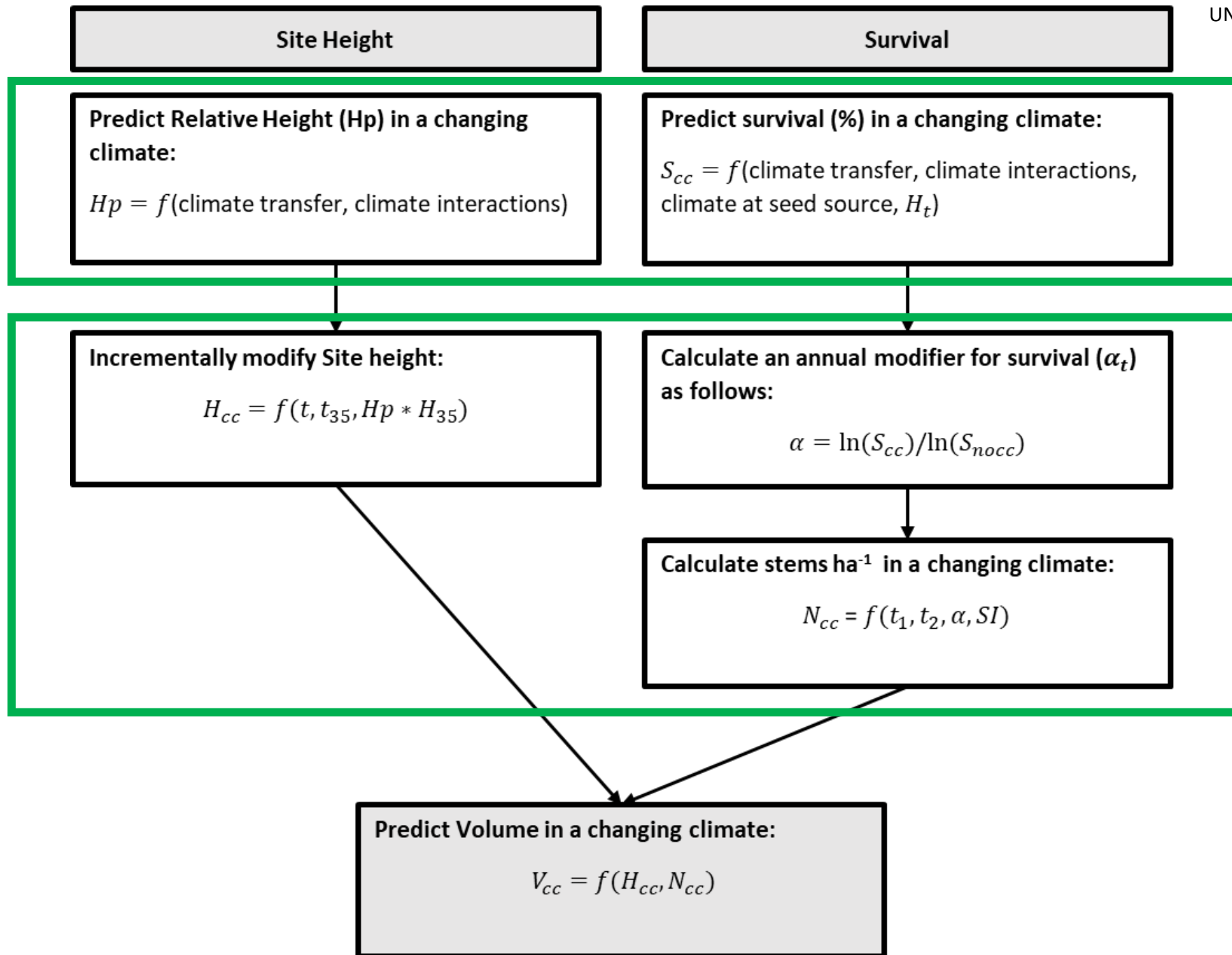
- **Modification of N (Stems/ha):**

- $\alpha = \ln(S_{cc}) / \ln(S_{nocc})$
- $N_{cc} = f(\alpha, N_{nocc})$



Transfer Functions

Application of modifiers

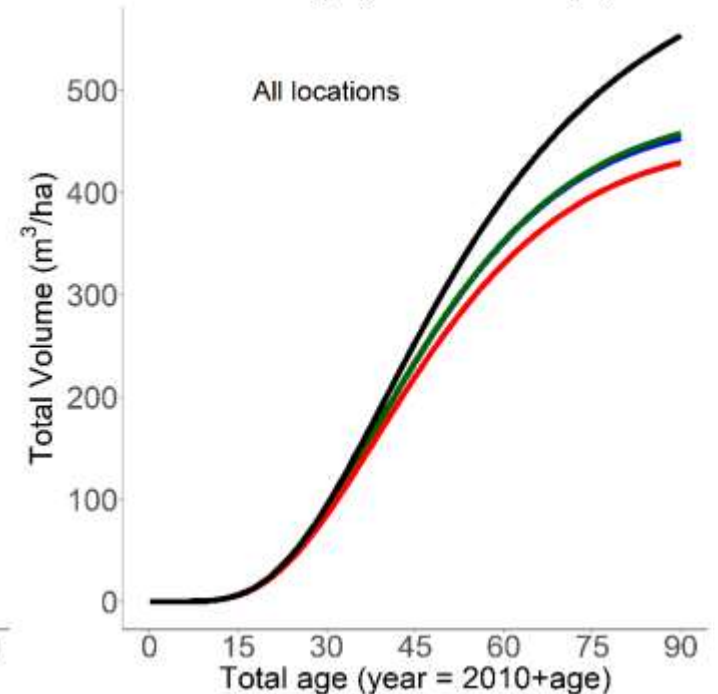
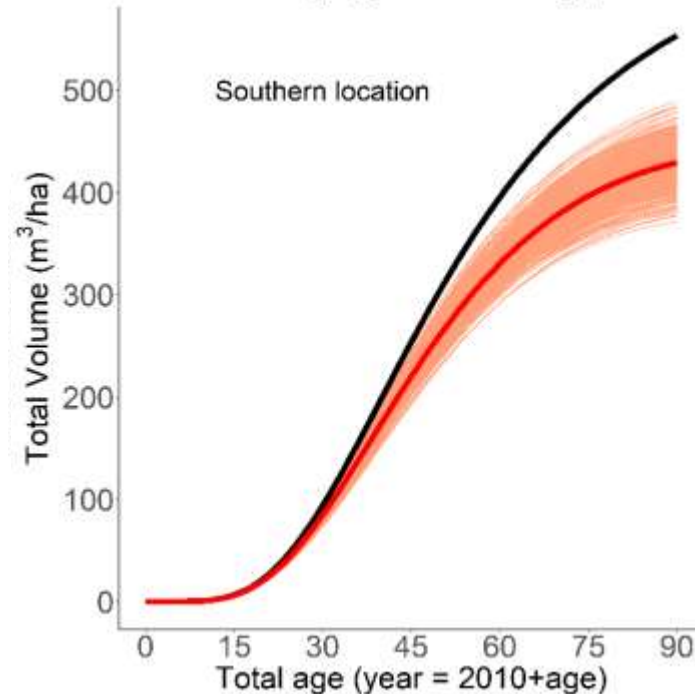
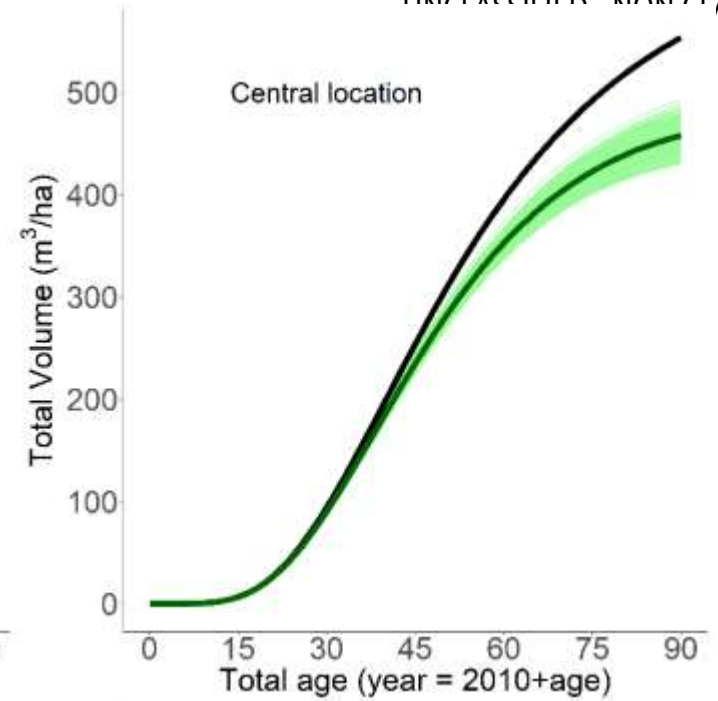
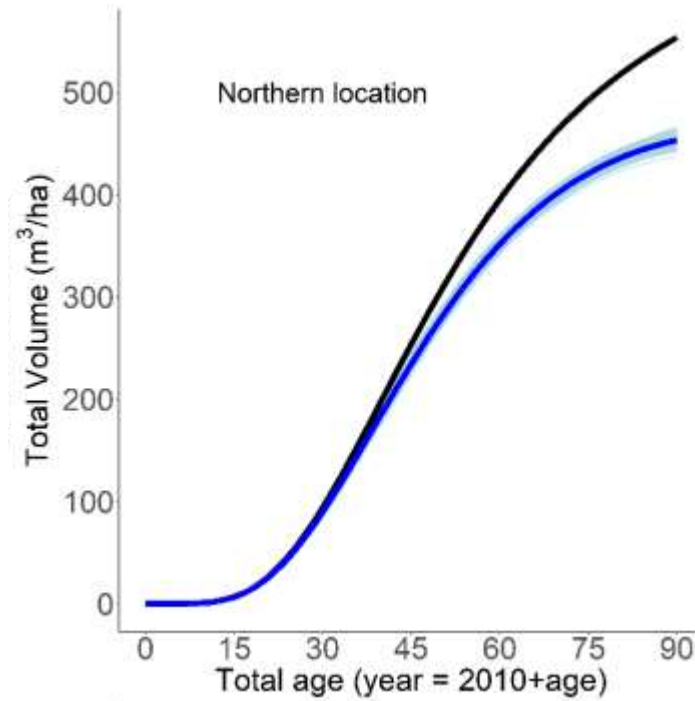


Simulations at three cutblocks (local seed)



Planted Pine
SPH = 1200
SI = 20 m
90 yr projection,
starting in 2010

17-22%
reduction in
stand volume



Limitations

- Provenance Trials \neq G&Y Plots
 - No buffers
 - Single planting density
 - < 18 trees / population
 - No crown measurements
 - Trials \ll Rotation Age
- Transfer functions for all species?
 - TFs for Generalists and Specialists?

Next Steps

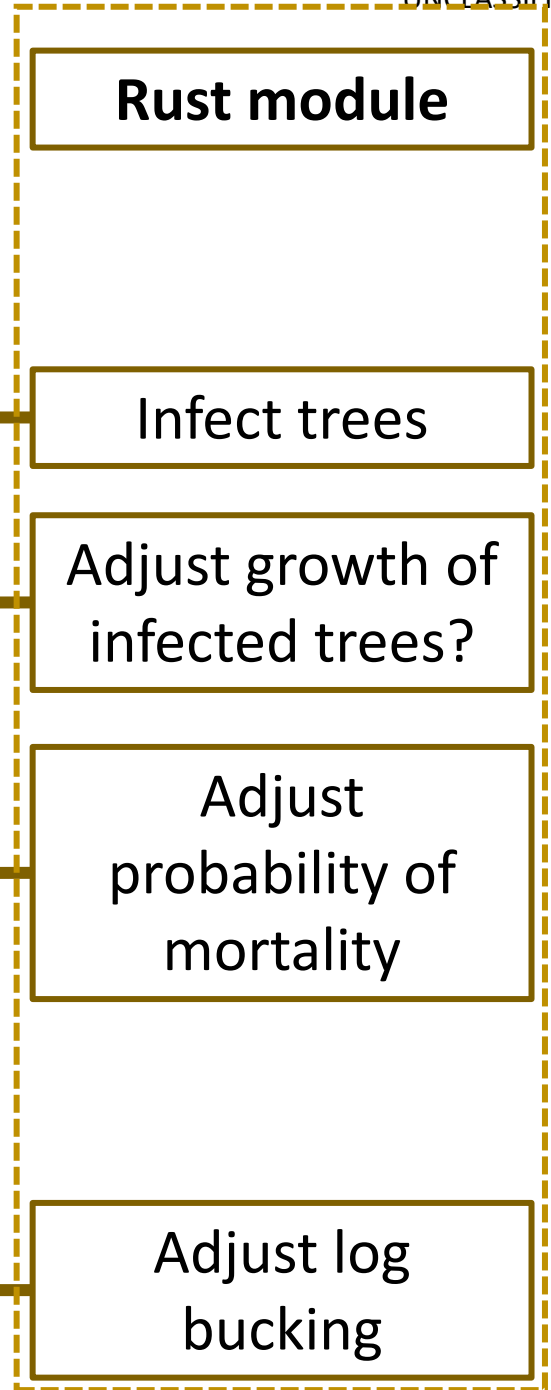
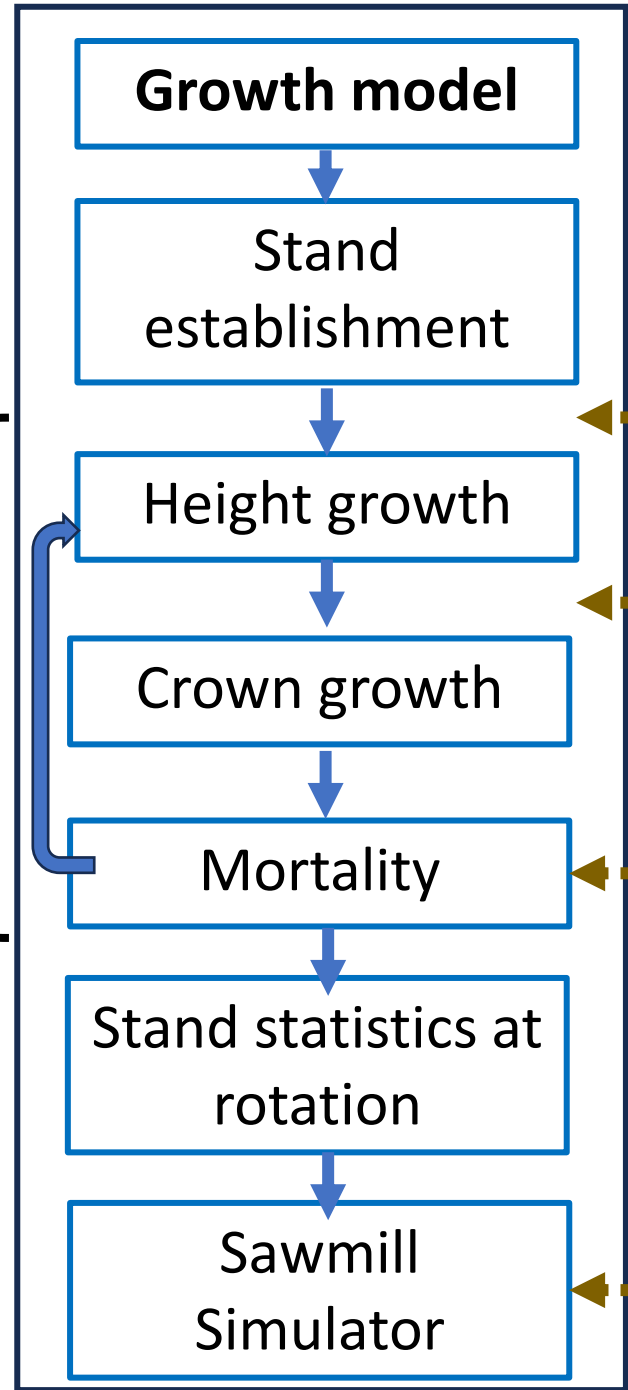
- Simulations with non-local seed
- Simulations with local/non-local seed mixtures
- Landscape-level analysis – harvest scheduler
- Adding the option to modify projections in TIPSYP???

Simulating the impacts of pine rusts





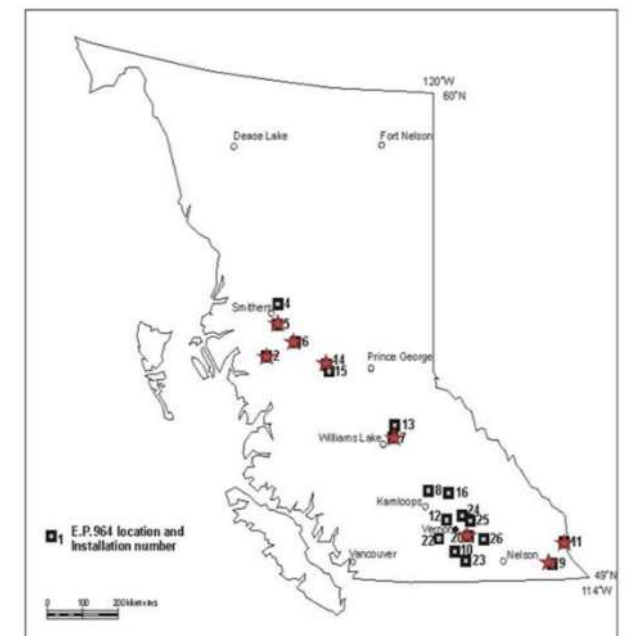
Annual
time-step



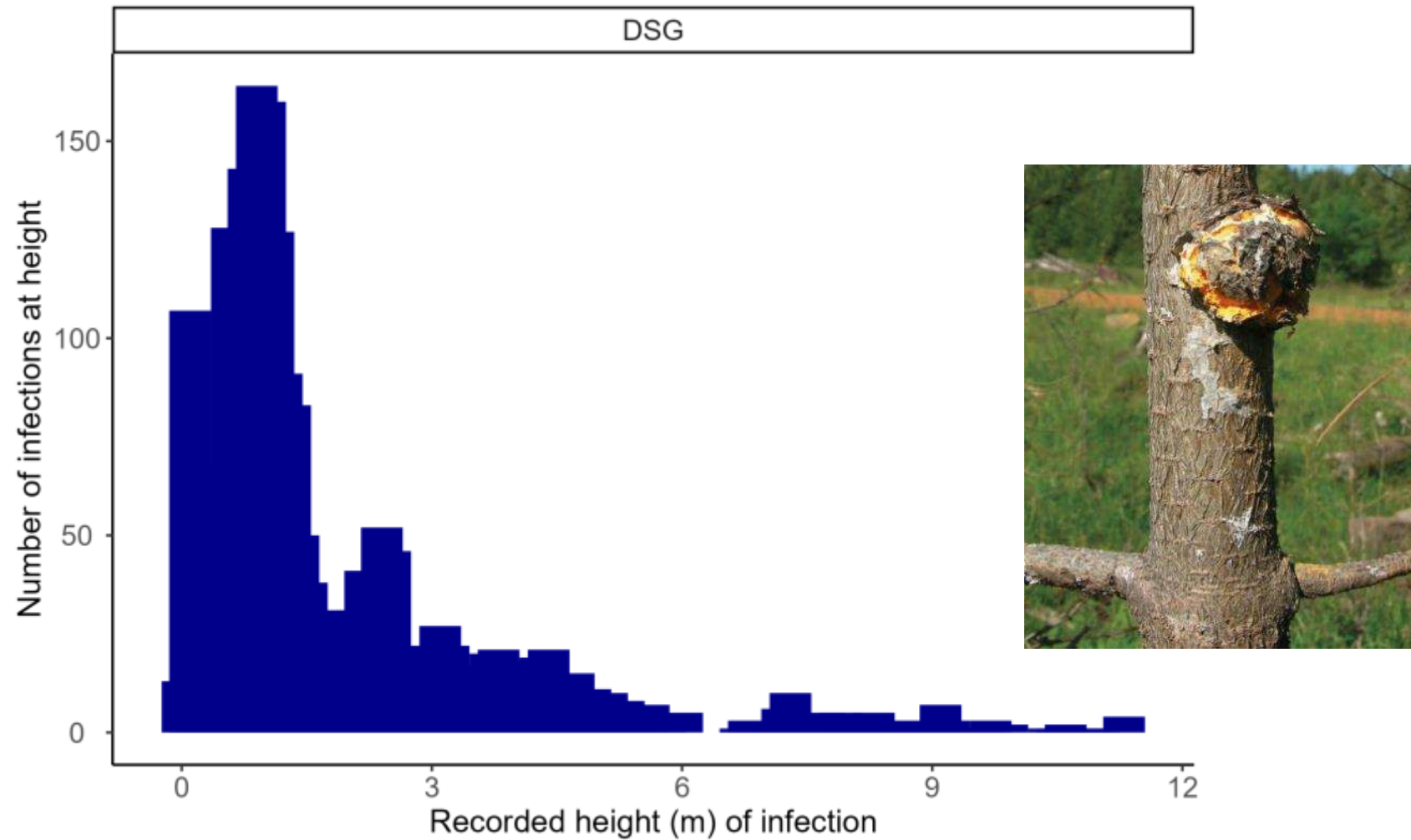
Data for model building

Main Data Sources:

- **EP 671 (1 installation; Est. 1968)**
 - 4 planting densities, each replicated in 6 plots (24 plots)
 - 64 trees/plot (+ buffers) measured at 8, 13, 18, 23, 28, 33, and 41 years
 - Rusts surveys performed in 1997, 2002, 2005, 2014
- **EP 964 (5 installations; Est. 1987-1991)**
 - 5 planting densities, each replicated in 3 plots
 - 64 trees/plot measured every 5 or 10 years, up to 25 years
 - Rust survey performed in 2016
- **Bednesti (1 installation; Est. 1988)**
 - Various planting densities + treatments over 30 plots
 - Most plots measured annually up to 30 years
 - Rusts surveyed in 2010, 2011, 2015, 2016, 2017

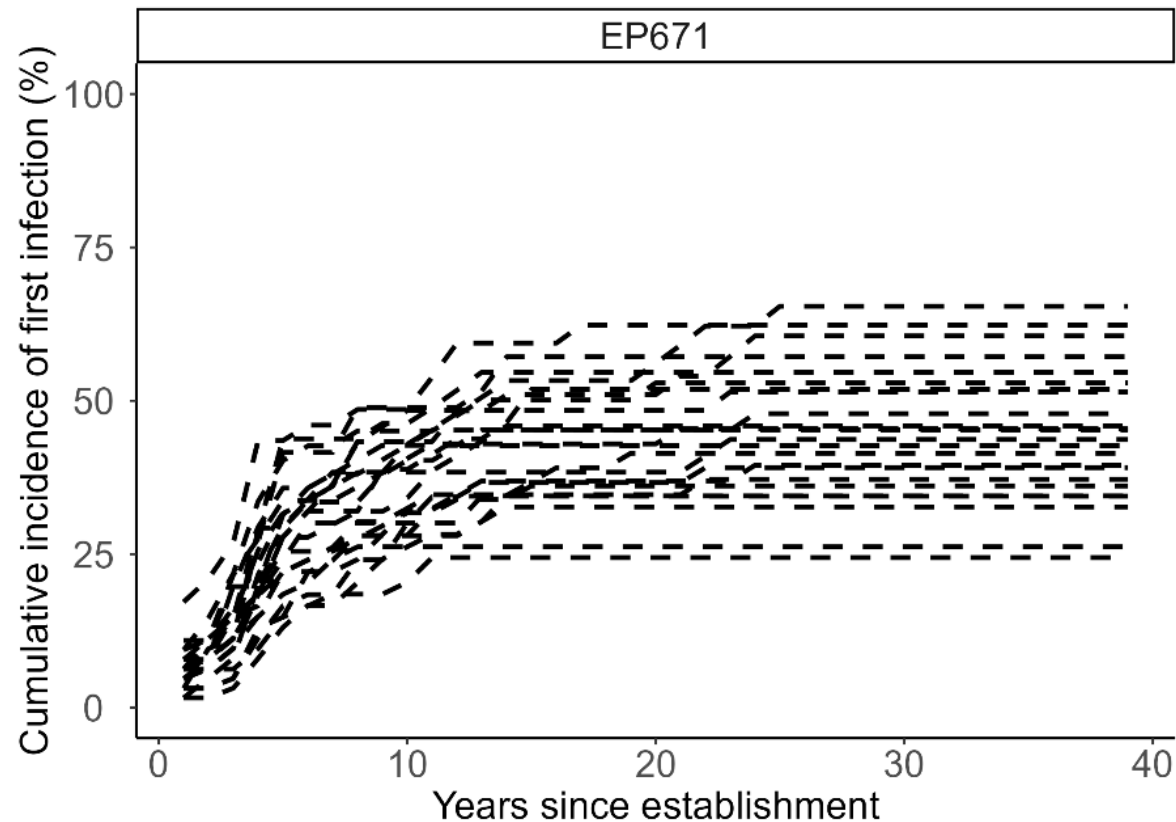


Heights of gall on main stem



Cumulative incidence of first infection

$$Cumulative_Inc = \sum_{t=1}^k Incidence\ Proportion$$



Rust module

Infect trees

Adjust growth of
infected trees?

Adjust
probability of
mortality

Adjust log
bucking

Cumulative incidence of first infection

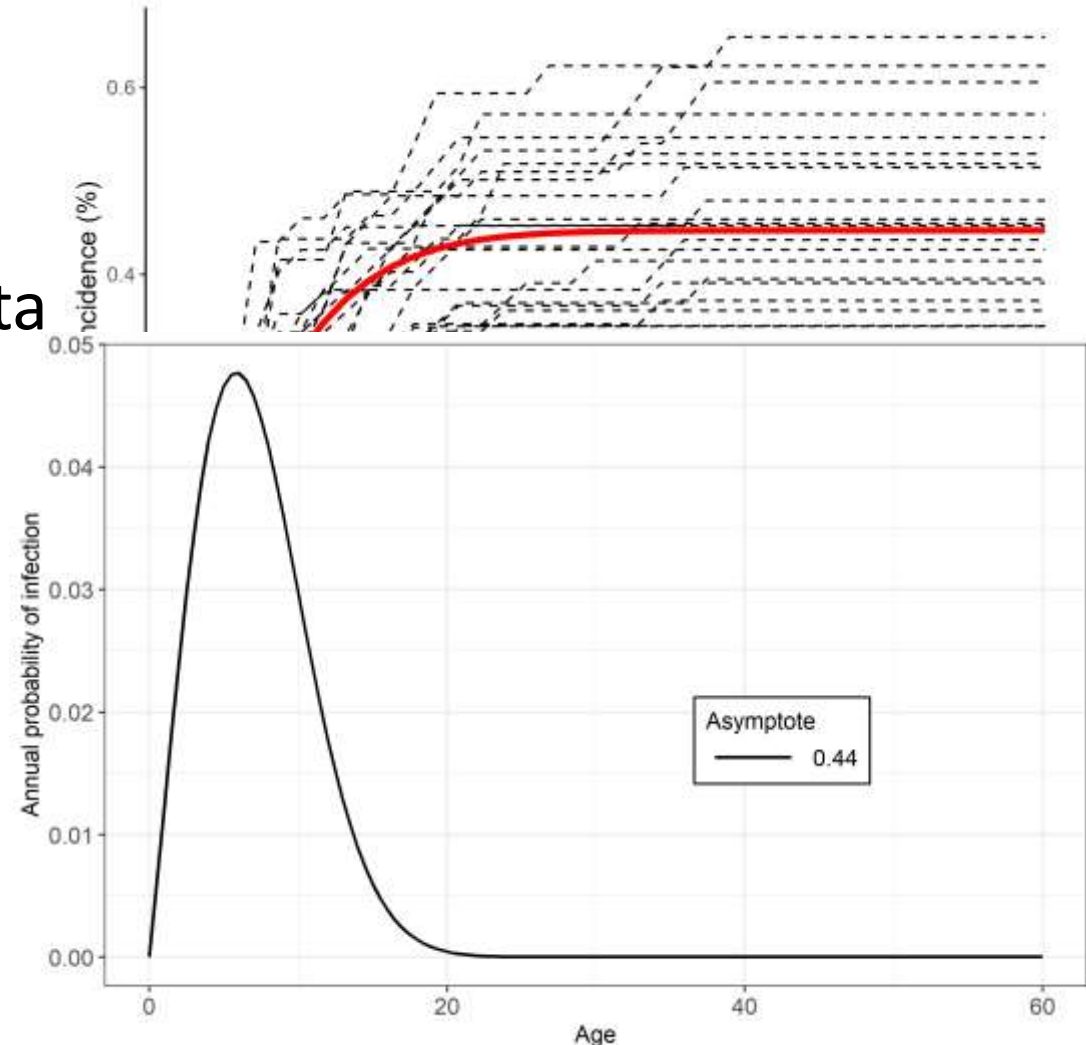
Weibull Growth Function

$$Cumulative_Inc = \beta_1 [1 - \exp(-\beta_2 t^{\beta_3})]$$

- Initialized using Free Growing Survey Data
- β_1 = initialization variable

Differential form (annual probability):

$$P_{inf} = \beta_1 \beta_2 \beta_3 t^{\beta_3 - 1} [\exp(-\beta_2 t^{\beta_3})]$$



Modified probability of infection

Two parameter Weibull:

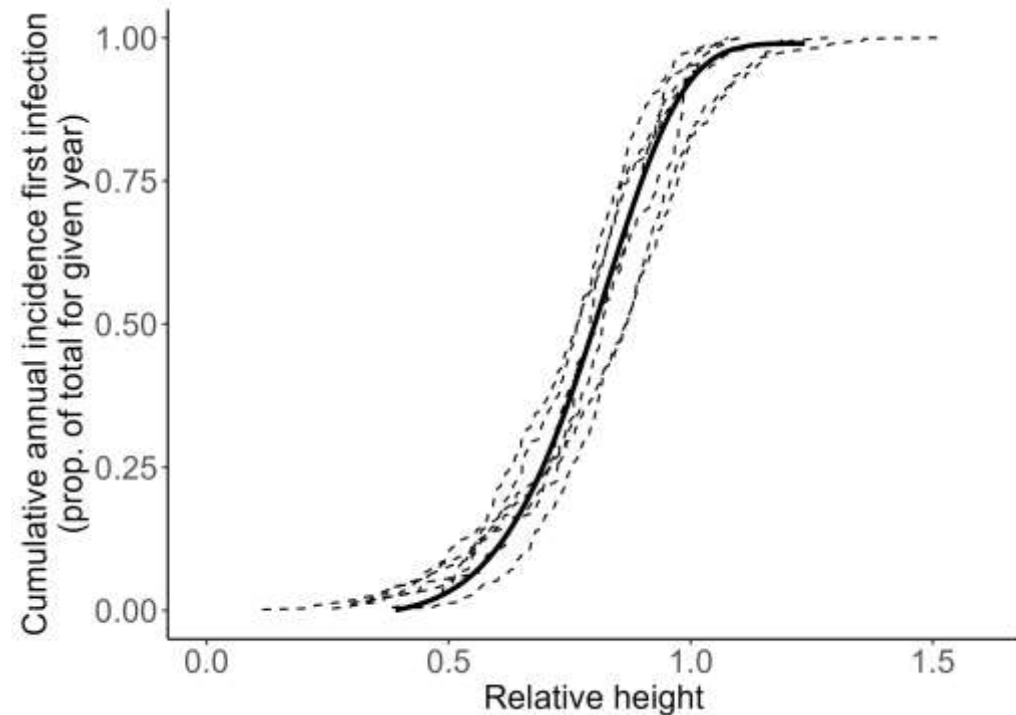
$$CAP_1 = 1 - \exp\left(-\left(\frac{RelH}{\alpha_1}\right)^{\alpha_2}\right)$$

Inverse transform:

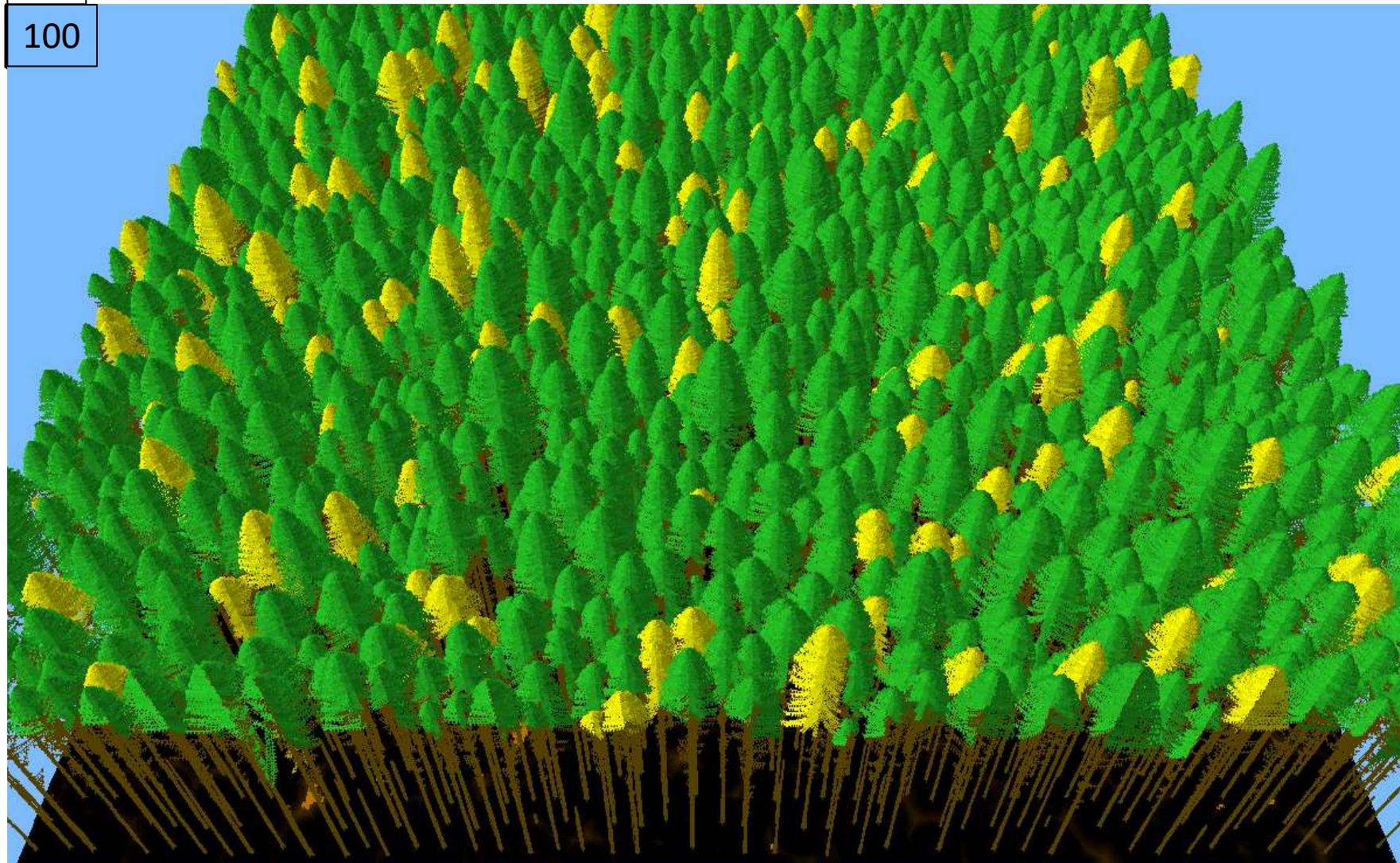
$$RelH_u = \alpha_1 [-\ln(1 - u)]^{1/\alpha_2}$$

(maps $U(0, 1)$ to Weibull distribution)

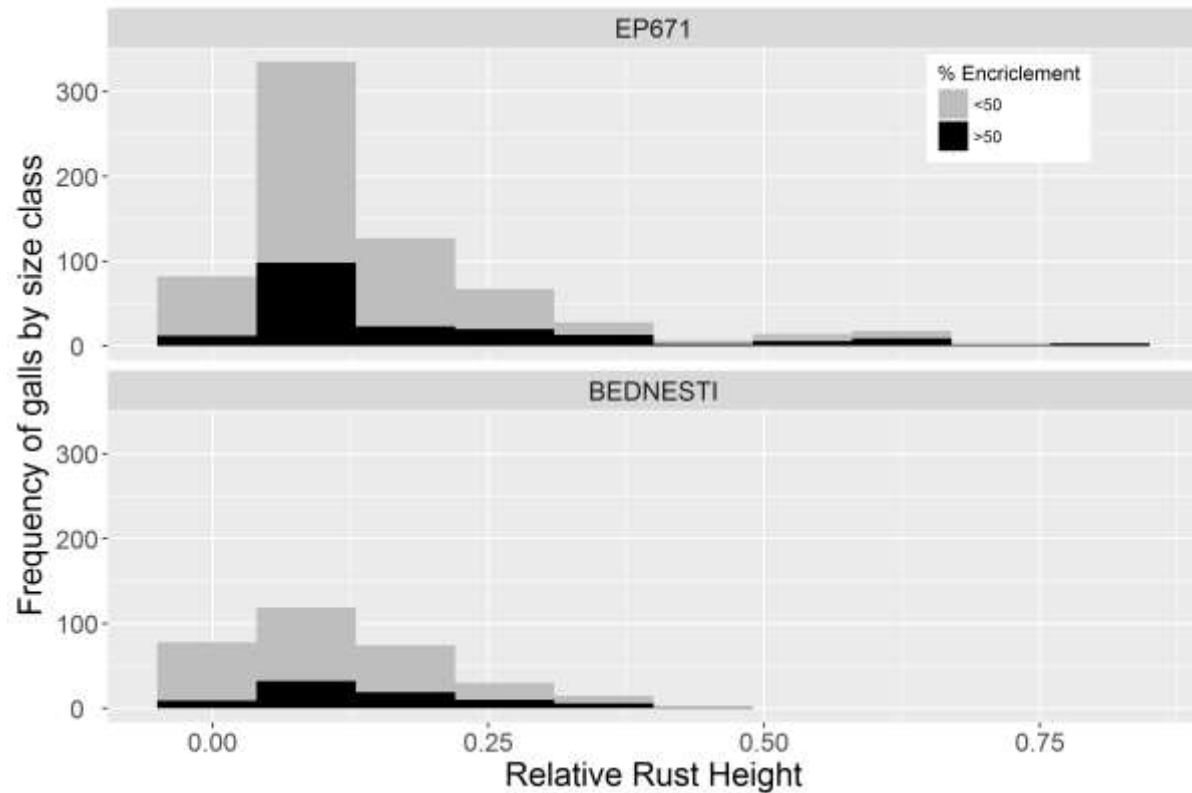
- Draw $u \sim U(0, 1)$, generate $RelH_u$
- Infect trees where: $RelH_i < RelH_u$
until we reach P_{inf} for that year



2500/ha planted, 20% incidence, standard juvenile mortality



Galls >50% encirclement



Rust module

Infect trees

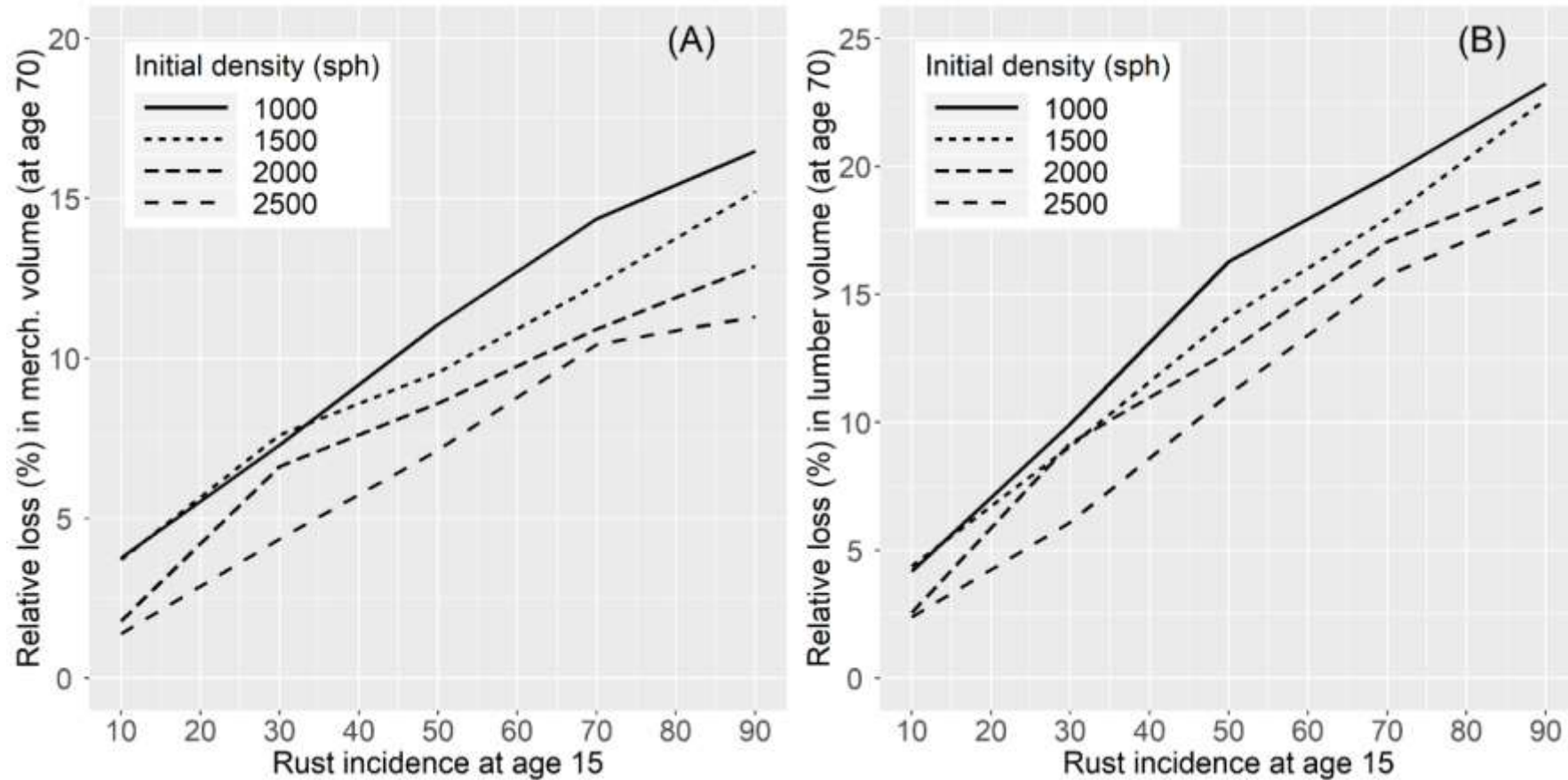
Adjust growth of
infected trees?

Adjust
probability of
mortality

Adjust log
bucking

- Is a Bernoulli random variable (either 0 or 1)
- If $> 50\%$ = 1, otherwise 0

Expected Loss in (A) Merch. Vol. and (B) Lumber



Questions?



Pedlar and McKenney (2017)

