

# PnET-Succession v4.0 Extension User Guide

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# 1 Introduction

This document describes the **PnET-Succession** extension for the LANDIS-II model. For information about the model and its core concepts including succession, see the [LANDIS-II Conceptual Model Description](#).

The PnET-Succession extension is based on the Biomass Succession extension of Sheller and Mladenoff (2004), embedding elements of the PnET-II ecophysiology model of Aber et al. (1995) to simulate growth as a competition for available light and water, replacing the existing competition for “growing space” algorithms. PnET (Photosynthesis and EvapoTranspiration) is a simple, lumped parameter model of carbon and water balances of forests (Aber and Federer 1992), built on two principal relationships: 1) maximum photo-synthetic rate is a function of foliar nitrogen concentration, and 2) stomatal conductance is a function of realized photosynthetic rate.

## 1.1 Major modifications made to PnET algorithms

Several modifications were made to PnET algorithms to make them tractable at landscape scales, primarily by broadening the scale of integration operations. (1) The timestep was broadened from the daily PnET timestep to monthly. (2) The number of sub-layers within a canopy layer was 50 in PnET, but is here set by the user (IMAX) to increase computational efficiency, where each sub-layer represents an even proportion of the total LAI within the layer. A greater number of subcanopy layers tightens the feedback between photosynthesis and water stress, but significantly increases computation time. (3) The original PnET adds foliage to successively deeper subcanopy layers until there is insufficient light to support photosynthesis. PnET-Succession allocates foliage in proportion to the active wood (xylem) that supports it. (4) Cohort biomass in PnET-Succession is used as a surrogate for tree height to simulate canopy layers, which are added when the variation in biomass among cohorts exceeds a user-defined amount. (5) Photosynthates in PnET-Succession are allocated to four pools (foliage, root, wood and non-structural carbon (reserves, NSC)). Net photosynthesis is initially allocated to the NSC pool, and then foliage allocation occurs, followed by allocation to root and wood pools such that the above- and below-ground biomass ratio is preserved. Maintenance respiration is then deducted from the NSC pool. Details of model structure and modifications can be found in De Bruijn et al. (2014).

## 1.2 Advantages and disadvantages of PnET-Succession compared to Biomass Succession

The goal for PnET-Succession was to make the simulation of growth and competition more mechanistic and more explicitly linked to fundamental drivers that are changing, such as climate and atmospheric composition (e.g., CO<sub>2</sub> and ozone). It is believed that this more mechanistic approach will be more robust for making projections under climate and other global changes (Gustafson 2013).

*Advantages of PnET-Succession compared to Biomass Succession*

- 1) PnET-Succession replaces the input parameters  $ANPP_{max}$  and  $B_{max}$  of LANDIS-II Biomass Succession with mechanistic and dynamic calculations of growth and senescence that depend on monthly climatic conditions and competition for resources. Establishment and growth are now emergent properties of the model and are explicitly linked to changing fundamental drivers such as climate and  $CO_2$  concentrations.
- 2) Dynamic calculations of LAI and photosynthesis allow cohorts to die prior to senescence, based on physiological constraints (too few carbon reserves). This can typically occur when carbon reserves production is insufficient to support growth due to shading, water competition, drought, diseases or pests. This allows more realistic simulation of cohort death in the course of stand development (i.e., mortality is highest in the younger cohorts), and a more realistic accounting of biomass accumulation. An added benefit is that the number of cohorts to be simulated is reduced.
- 3) PnET-Succession allows a more explicit simulation of species' survival strategies, by implementing a dynamic competition for light and water. For example, one can parameterize species or species-group combinations of respiration losses and water use efficiency to implement competition advantages or disadvantages for particular species in sites that are dry or shaded due to competing vegetation. PnET-Succession also allows simulation of waterlogging effects on photosynthesis.

#### *Disadvantages of PnET-Succession compared to Biomass Succession*

- 1) PnET-Succession requires more parameters, which adds to uncertainty and increases the parameter burden when using the model. However, uncertainty may be no higher than when making *ad hoc* assumptions for other succession extensions about how novel conditions will affect modeled processes.
- 2) Runtimes tend to be longer because there are more calculations for each cohort. However, the number of cohorts may be less because many cohorts dies of stress or senesce prior to reaching longevity age, greatly reducing the number of cohorts that must be simulated.

### 1.3 What's new in Version 4.0 (December 2020)

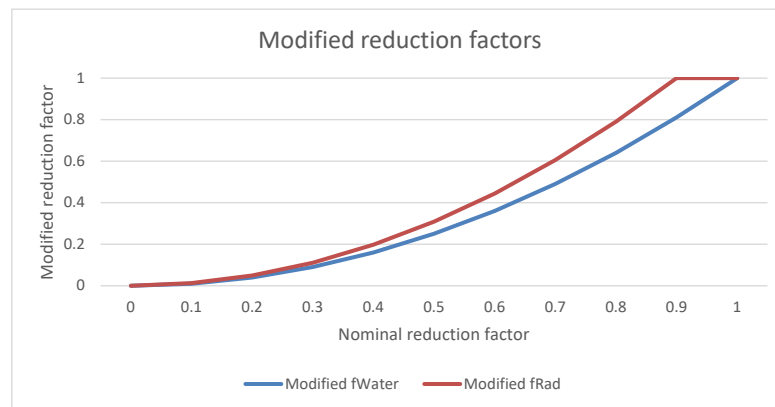
Version 4.0 modifies the establishment algorithms to improve control of establishment and to reduce the proliferation of cohorts that have little chance of surviving, which can lengthen run times and consume computer memory. The EstMoist and EstRad parameters previously had no effect when a reduction factor equaled (or approached) 1.0, providing limited control over establishment. These parameters are no longer compatible with those of prior versions, but all other inputs from prior versions remain compatible.

Version 4.0 adds optional capabilities to simulate hydrologically waterlogged soils, specifically forested wetlands and forests on permafrost. Forested wetlands are poorly simulated in most forest landscape models, so this is an important advance.

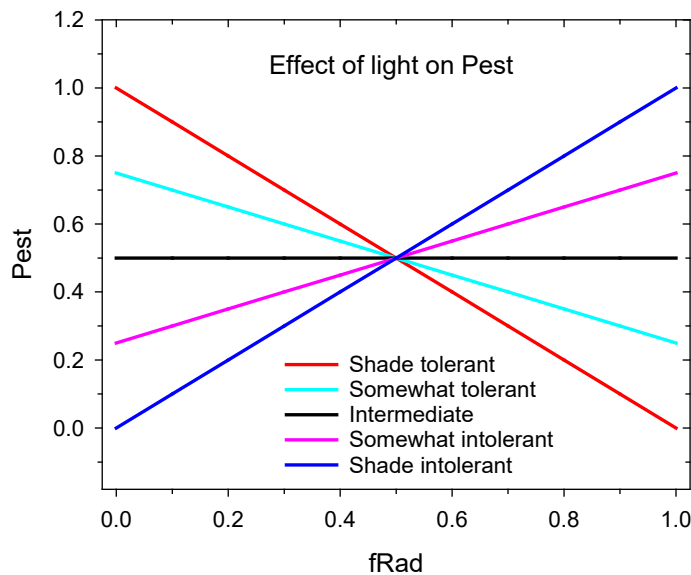


### 1.3.1 Added Features

- The EstMod parameters have been repurposed to make the light and water effects on establishment to be independently scalable. EstRad now specifies the fRad above which the light environment is optimal for establishment, and EstWater specifies the fWater above which the soil moisture is optimal for establishment. Although fWater often takes values of 1.0, fRad almost never does, and these parameters allow the user to scale them to have equal effect, if desired. One approach to setting these parameters is to make them equal to the highest expected value for each reduction factor (typically the maximum seen in calibration runs when there is no competition for light), which is often 1 for fWater, and lower for fRad. Both reduction factors become scaled equally across the range of their expected values when computing the probability of establishment (Pest). The figure below represents the impact of the EstMod parameters with value of EstRad = 0.9 and EstWater = 1.0.



- OPTIONAL:** The effect of light on establishment (Pest) can be set to no longer be directly proportional to fRad. Instead, Pest is highest for species with high HalfSat (pioneers) when light is high, and highest for shade-tolerant species when light is low. Pest is intermediate for species with mid-range HalfSat in all light environments. The effect of water on Pest remains proportional to fWater. This modification greatly improves the ability of the model to establish species where they are most likely to thrive and to not establish them where they will not be competitive, reducing model runtimes.



- Parameter MaxPest allows global tuning of absolute establishment probabilities. The EstMod parameters equalize the effect of light and water and links them to their tolerance of each factor, and MaxPest adjusts the absolute values of all Pest computations up or down equally to control overall cohort proliferation.
- Cohorts are now killed if their biomass drops below initial biomass at the end of a year. Such cohorts almost always eventually die, but they may otherwise take 40+ years to die because their maintenance respiration costs are very low.
- To optionally simulate soil ice dynamics (specifically, permafrost), the rooting depth and leakage parameters are now dynamic at a monthly time step, determined by ice depth as driven by the temperature inputs. Rooting depth in this case is understood to represent the mean depth to the ice layer (0 °C isotherm) in the month, constrained to not exceed the user-defined RootingDepth. When the ice layer retreats below the rooting depth, we assume that the water column within the root zone is increasingly able to drain, and the leakage fraction parameter is therefore increased linearly from 0.0 (when ice occurs within the root zone) to 1.0 when the ice layer is at least a user-specified depth (LeakageFrostDepth) below the root zone. Thus, the presence of permafrost varies the water potential within the rooting zone according to temperature dynamics. See Gustafson et al (in press) for details.

### 1.3.2 Bug Fixes

- Potential evaporation calculation was modified for consistency in units, using a different formulation (see 2.4.2.2)
- Corrected rounding error that sometimes allowed soil water to be negative under extreme dry conditions.

- Modified output metadata to improve compatibility with the LandViz visualization tool.

## 1.4 References

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## 1.6 Release History

### 1.6.1 Major Releases

#### 1.6.1.1 Version 3.0 (June 2018)

##### *1.6.1.1.1 Added Features*

- Latitude can now be used as an ecoregion parameter (or as a global parameter as before), allowing for large study areas with spatially dispersed ecoregions. Latitude is used to compute day length available for photosynthesis.
- PnET-Succession v2.0 (and earlier) used equations from early versions of PnET-II to compute the CO<sub>2</sub> enrichment effect (DelAmax). The most recent version of PnET-II uses an equation from Franks et al. (2013), which moderates the CO<sub>2</sub> effect, especially at high CO<sub>2</sub> concentrations. We now use a modified version of the Franks equation that uses internal leaf CO<sub>2</sub> concentrations rather than external leaf CO<sub>2</sub> concentrations (Gustafson et al 2018). The Franks equation lessens the CO<sub>2</sub> enhancement effect, especially at high CO<sub>2</sub> concentrations.
- Now PnET-Succession computes a variable (CiModifier) used to modify internal leaf CO<sub>2</sub> concentration (Ci), reflecting reduced conductance as a function of drought stress (fWater), ozone dose, and species ozone tolerance. This is used to reflect stomatal closure caused by water stress, which is modified by stomatal sluggishness (inability to close completely) induced by elevated ozone, modeling its interaction with conductance of CO<sub>2</sub> and water. Absorption of CO<sub>2</sub>, O<sub>3</sub>, and transpiration of water are all reduced by CiModifier.
- Modified the canopy layering algorithm so that all cohort canopy sublayers are assigned to only one main canopy layer (i.e., a cohort cannot span multiple main canopy layers).
- Provided a parameter (CO2HalfSatEff) to optionally make HalfSat dynamic as a function of CO<sub>2</sub> concentration. This effect can be turned off by setting to zero.
- Provided parameters (FolNInt, FolNSlope) to optionally make FolN dynamic as a function of light (fRad), allowing photosynthetic capability (Amax) to vary vertically through the canopy (by canopy sublayer) and in response to cohort release or overtopping. This dynamic effect can be turned off by setting these to 1.0 and 0.0, respectively.
- Added optional output variables to the PnETOutputSites file with species-specific amounts of dead wood (WoodySenescence) and foliage (FoliageSenescence) while

tracked internally before, were not available as outputs from the respective dead biomass pools. This output is useful for more landscape carbon accounting.

- Dropped the WUEConst species parameter. Water use efficiency is now calculated directly from fluxes of water from leaves ( $J_{H_2O}$ ).
- Activated the H2 species parameter that allows simulation of waterlogging effects on photosynthesis.
- Added the ability to optionally include ozone effects on photosynthesis. These functions are activated only when the climate input file contains a field with monthly cumulative ozone concentrations.
- Added the O3GrowthSens parameter to scale the computation of the species-specific ozone effect that reflects the damaging effects of ozone on photosynthetic tissue. This is used to specify species sensitivity to ozone damage of tissues, and is only needed when ozone data are provided in the climate file.
- Added the O3StomataSens parameter to reflect species differences in stomatal sluggishness when exposed to ozone. This categorical class impacts how CiModifier is calculated, and is only needed when ozone data are provided in the climate file.
- Litter decomposition is now computed once per year, including during initial spin-up.
- Added DisturbanceReductions that allow disturbances to impact the existing dead wood and litter pools. Other disturbance impacts on biomass transferring from live to dead pools that used to be included in the AgeOnlyDisturbances file have been moved to the DisturbanceReductions file. Added section to the Users Guide to explain the DisturbanceReductions.

#### ***1.6.1.1.2 Bug Fixes***

- Bug fix to ensure all foliage and NSC are lost when a cohort dies, even though it is not removed until the end of the succession timestep. This step makes the cohorts functionally dead the first year the NSCFrac drops below 0.01.

### **1.6.1.2 Version 2.0 (October 2016)**

#### ***1.6.1.2.1 Added Features***

- New generic parameter: PrecipEvents. Divides incoming monthly precipitation into discrete events within the month ( $n = \text{PrecipEvents}$ ) and applies each portion randomly during the sequence of processing canopy sublayers. This prevents large cohorts from consuming a disproportionate share of the available water in a given month.
- New generic parameter: Wythers. Option to apply the foliar respiration modification to increased temperature as described by Wythers et al (2013).

- New generic parameter: DTEMP. Option to apply the temperature reduction factor (DTEMP) of PnET-II rather than the original PnET-Succession v1.2 temperature reduction factor. The v1.2 temperature reduction factor does not explicitly penalize photosynthesis at temperatures above PsnTOpt.
- New ecoregion parameter: SnowSublimFrac. Snowpack is reduced by this amount prior to snowmelt, representing sublimation and meltwater runoff that does not enter the soil.
- New output options for woody senescence and foliage senescence by species.

#### ***1.6.1.2.2 Bug fixes:***

- A bug in the calculation of transpiration was fixed.
- A bug that caused the decomposition of dead pools to not be simulated during spin-up in prior versions was fixed.
- A bug in the calculation of runoff was fixed.
- Biomass values provided to disturbance extensions in prior versions were the sum of above- and below-ground woody biomass, but no foliar biomass. Version 2.0 includes aboveground and foliar biomass to be consistent with other Biomass Succession extensions and is therefore more compatible with biomass disturbance extensions. Specific biomass pools can be now output as maps and total pool sizes using the Output-PnET extension (Section 10).
- Defoliation (by an external disturbance extension) is now applied during June (previously it was January when deciduous species had no foliage).
- A bug in the processing of cohorts killed by disturbance was fixed. The bug prevented disturbances from recording the cohorts being removed.
- A bug in the calculation of snow melt was fixed. The bug caused all snow pack to melt at the same time. The rate of snowmelt was changed to 2.74 mm/°C/day (see 2.4.2.1).
- When snowpack is present, surface PAR is set to 0 which eliminates soil water evaporation under snow (though sublimation of snow occurs instead).

An Excel worksheet [PnET-Succession function worksheet.xlsx] is available from (<http://www.landis-ii.org/extensions/pnet-succession>) that can be used to better understand how selected input parameters affect state variable computations.

#### **1.6.1.3 Version 1.0 (2014)**

Original released version

## 1.6.2 Minor Releases

### 1.6.2.1 Version 4.0.1 (January 2021)

Fixed a bug that caused initial cohorts with age 1 to not successfully initialize through spin-up. This bug was introduced in v4.0 and only occurred when cohorts in the initial community file had an age of 1 and no initial biomass provided.

### 1.6.2.2 Version 3.4 (December 2019)

Version 3.4 added several new features and corrected some important bugs. Nearly all of the added features are optional, with the intention of maintaining compatibility with inputs that worked in the previous version.

#### ***1.6.2.2.1 Added Features***

- Optional ability to use the Climate Library to provide climate data.
- New optional PsnTMax species parameter to set the upper temperature limit for photosynthesis. Previously, this was estimated from PsnTMin and PsnTOpt, assuming a symmetrical relationship so that  $(PsnTOpt - PsnTMin) = (PsnTMax - PsnTOpt)$ .
- New optional CohortBinSize parameter to aggregate cohort age bins to be larger than the extension timestep.
- Optional ability to supply cohort initial biomass in the initial communities file.
- Optional reading of initial litter and dead wood maps.
- New optional RunoffFrac ecoregion parameter to allow prevention of some water runoff.
- New optional LeafOnMinT species parameter to allow separate control of the beginning and end of the growing season, apart from the control of photosynthesis rate at low temperatures (PsnTMin).
- New optional ColdTol species parameter to identify the minimum temperature for survival of the species cohorts.
- New WinterStd ecoregion parameter to estimate the extreme cold temperature from the monthly average TMin.
- Added EstablishmentTable output option.
- Added MortalityTable output option.
- Additional information in output files.
- Cohorts can re-flush foliage following defoliation caused by a disturbance extension. Refoliation occurs at 70% of ideal foliage, for deciduous ( $TOfol = 1$ ) cohorts that experience  $> 60\%$  defoliation in a given year. Refoliation has additional cost to NSC, and cohorts that do not re-leaf still experience additional NSC losses.

#### ***1.6.2.2.2 Bug Fixes:***

- Correction to the initialization of foliage for initial communities that are replicated in the initial community map. Failure to initialize foliage correctly caused cohorts to die immediately following spin-up, but only on sites that were duplicated in the initial communities map, and were not included as PnETOutputSites.

- Correction to the accumulation of cohort biomass when combining multiple cohorts less than the timestep (or CohortBinSize) age.
- Correction to the calculation of GrossPsn that resulted in cases of transpiration exceeding soil water.
- Correction to the DTemp response calculation that resulted in values <0.

#### 1.6.2.3 Version 3.3 (Internal Release)

Added support for providing initial cohort biomass (no spin-up)

#### 1.6.2.4 Version 3.2 (Internal Release)

Added support for Climate Library.

#### 1.6.2.5 Version 3.1 (September 2018)

Version 3.1 is compatible with the LANDIS-II core v7.

Added dynamic foliage responses to light (fRad) through dynamic foliar nitrogen (FolN) and foliage fraction (FracFol). See sections 8.25 and 8.26.

#### 1.6.2.6 Version 2.1.1 (October 2017)

This release incorporates a change to the Biomass Cohort Library to maintain compatibility with other extensions that use the same library (all extensions that use cohorts with biomass attributes). The edit to the Biomass Cohort Library enabled the proper tracking of dead biomass (additions to the dead pools) when partial cohort removals occurred.

This version also adds compatibility with the Metadata Library that supports output visualization using the VizTool.

#### 1.6.2.7 Version 2.1 (May 2017)

Rename [SurfaceRunoff] to [PrecLoss] in the Site Output file to distinguish between water lost due to soil saturation (SurfaceRunOff) and water lost due to other factors (PrecLoss; e.g., slope, impervious surface). Add tracking of PrecLoss variable.

Rename [Layer] to [TopLayer] in the Cohort Output file to denote that the value reported is the highest layer in which the cohort appears. The top canopy layer has the highest layer value.

The allocation of precipitation events to subcanopy layers has been adjusted so that the precip events are randomly assigned to layers, but not constrained to a single event per layer. This can result in multiple precip events occurring (with their associated runoff, interception, leakage, etc.) for a given layer, especially when the number of precipitation events is greater than the number of subcanopy layers on a site. This resolves a discontinuity in the water cycle when the number of cohorts was low relative to the number of precip events.

##### ***1.6.2.7.1 Bug Fixes:***

- An important bug related to dispersal was fixed in this version. Previously, the age of the **youngest** cohort of a species was used to determine if a mature cohort



was present on a site for seeding purposes. The test should use the age of the **oldest** cohort of a species to check for maturity and determine sources of seed for dispersal. This error has been corrected.

- Fixed the DiscreteUniformRandom function to be inclusive of the maximum value. Previous implementation may have been slightly biased in the shuffling of subcanopy layers.

## 2 PnET-Succession

The PnET-Succession Extension generally follows the methods of the Biomass Succession Extension: Age cohorts reproduce, grow (add biomass), age, and die. The PnET-Succession Extension replaces the simple growth and competition algorithms from the Biomass Succession Extension with the photosynthesis and respiration equations from PnET-II to simulate growth of specific cohort biomass components (root, foliage, wood, non-structural carbon) as a competition for water and light.

PnET-Succession simulates the competition of cohorts for water and light as a function of photosynthetic processes. Competition for water is simulated on each site (grid cell) through a dynamic soil water balance that receives precipitation and loses water as runoff, interception, percolation out of the rooting zone and consumption by cohorts through transpiration. Competition for light is modeled by tracking solar radiation through canopy layers (related to cohort age) according to a standard Lambert-Beer formula. PnET-Succession requires average monthly temperature, precipitation, photosynthetically active radiation and atmospheric CO<sub>2</sub> concentration as inputs.

Because monthly climate data are provided as an input to the extension, species establishment probability is also calculated at each time step as a function of the climate conditions during the time step.

The PnET-Succession Extension also changes the calculation of shade. LAI is estimated for multiple canopy layers, and available light is computed for each layer, including the sub-canopy (i.e., ground).

The PnET-Succession Extension tracks biomass in four live pools (foliage, roots, wood and non-structural (C reserves)) and two dead pools (woody and leaf litter). For disturbance extensions that request “biomass” from the succession extension, PnET-Succession passes the sum of wood+foliage.

### 2.1 Initializing Biomass

At the beginning of a scenario, the initial communities begin with appropriate living and dead biomass values estimated for each site. Users have the option of providing initial aboveground live biomass values for each cohort, or allowing PnET-Succession to estimate the biomass. The required format for initial communities with biomass values is consistent with the text and map formats produced by the Biomass Community Output extension. When initial biomass values are not provided, the PnET-Succession extension uses its growth algorithms to iterate the number of time steps equal to the maximum cohort age for each site. Beginning at time (t - oldest cohort age), cohorts are added at each time step corresponding to the time when the existing cohorts were established. Thus, each cohort undergoes growth and mortality for the number of years equal to its current age, and its initial biomass value reflects competition among cohorts. Note: this is a computationally intensive process that may require significant time for complex initial landscapes. Additionally, climate data are required back to t - oldest cohort age. To facilitate climatic input in years where weather records do not exist, it is possible to supply mean monthly climate data for a range of years (see section 6.1.2)

This biomass initialization does not account for disturbances that would likely happen prior to initialization and therefore tends to overestimate initial live biomass and underestimate initial dead biomass. Because PnET-Succession gives bigger cohorts priority access to light and water, older cohorts may have an advantage during spin-up. Furthermore, some cohorts may not survive spin-up.

## 2.2 Cohort Competition

Biomass growth is driven by photosynthesis, which depends on light, soil moisture and foliage biomass. Multiplicative reduction factors are applied to gross photosynthesis to account for water stress, suboptimal radiation, vapor pressure deficit, and temperature. A similar growth enhancement factor (DelAmax) is applied for CO<sub>2</sub> concentrations above 350 ppm.

### 2.2.1 Light

The reduction of radiation intensity through the canopy is estimated using an exponential decrease function (i.e., Beer-Lambert law), where incoming radiation drives photosynthetic activity (Aber and Federer, 1992). A laboratory-derived relationship between foliar nitrogen concentration and assimilation rates under optimal growth conditions is used to estimate potential gross photosynthesis.

PnET-Succession assumes that LAI and biomass are spatially homogeneous on a site (i.e., cell). PnET-Succession defines canopy layers according to biomass, with high biomass cohorts achieving dominance with regard to access to light. Because senescing cohorts are more likely to lose biomass due to death of suppressed individuals or branches breaking off rather than due to the top breaking (i.e. they lose biomass without losing height), maximum lifetime biomass of a tree-species cohort is used as a proxy for tree height. Note that changes have been made to the model described paper in Ecological Modeling (De Bruijn et al 2014). We no longer set (arbitrary) age limits on the development of canopy layers. Rather, each cohort is divided into a number IMAX (=5 by default, set in the GenericParameters file) of canopy sublayers of equal biomass (Figure 1, left). The cohort sublayers are overlaid and the model iteratively clusters sublayers into canopy layers until the deviation of the newly formed canopy boundaries (i.e. LyrMax -LyrAv) decreases below a user defined parameter MaxDevLyrAv (Figure 1, right). This process produces boundaries of canopy layers such that the variation of subcanopy boundaries within the canopy layers is minimized and the variance of subcanopy boundary between canopy layers is maximized. MaxDevLyrAv is calibrated by the user to produce the maximum number of canopy layers expected in the system, or to control how long it takes for multiple canopy layers to form. The maximum number of canopy layers can further be limited with the generic MaxCanopyLayers parameter (section 7.4). Subcanopy layers within a canopy layer have equal access to the light reaching the canopy layer, and the light passing through a canopy layer is a function of the LAI and extinction coefficient of the cohorts making up the canopy layer. Light stress for a cohort is calculated by:  $fRad = Radiation / (Radiation + HalfSat)$ .

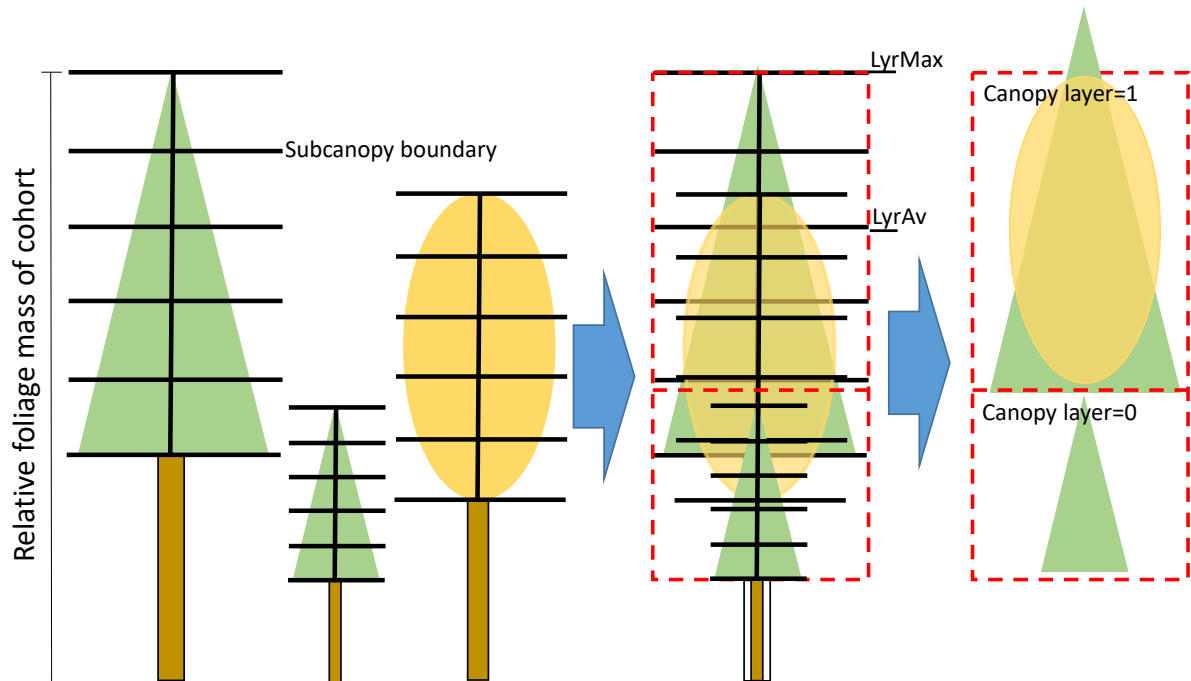


Figure 1. Canopy layer assignment. Each tree represents the foliage of a species-cohort on a site, and the solid horizontal lines are subcanopy boundaries (IMAX=5). Dotted lines are computed canopy layers that minimize the variance of foliage biomass within canopy layers and maximize the variance between canopy layers. All cohort subcanopy layers are assigned to the single canopy layer in which most of the cohort's subcanopy layers occur. Note that the diagram shows canopy shape only to enhance interpretation; in the model, subcanopy layers represent cohort foliage across the site, not tree crowns, and they all have equal biomass.

### 2.2.2 Water

Soil water is calculated in a bulk-hydrology model that updates soil water depending on precipitation, evaporation, soil drainage and consumption by the trees (Figure 2).

#### 2.2.2.1 Water In

Sources of bulk soil water (*soil\_water* in mm) are precipitation (*P* in mm/month) and melting water (*snow\_melt* in mm/mo). Incoming precipitation is intercepted by existing foliage at a rate controlled by a user parameter (*PrecipIntConst*), which defines the proportion of precipitation intercepted for each unit of leaf area (*LAI*).

$$Interception = P \times e^{(-PrecipIntConst * LAI)}$$

Intercepted precipitation is assumed to evaporate from the leaf surface and does not enter the soil. Sites with no live cohorts have no precipitation interception. When average temperature is below freezing, precipitation (snow) is not subject to interception and is allocated to *snow\_pack* (mm), where it remains until air temperature induces melting. Sublimation of snow is modeled as a direct reduction of snow pack prior to melting at a rate set by the ecoregion parameter *SnowSublimFrac* (default is

0.15 [Hood et al. 1999]). At above-freezing temperatures, snow melts at a rate of 2.74 mm/°C/day (USDA NRCS 2004). Snow melt is not subject to interception by foliage.

The combination of non-intercepted precipitation and snow melt define the potential incoming water. The incoming precipitation is divided into discrete precipitation events, with the number of events within a month set by the PrecipEvents ecoregion parameter. Precipitation from individual events is randomly assigned to individual subcanopy layers, with some layers potentially receiving multiple events and others receiving none. This random assignment ensures that all layers have equal priority to the incoming water. The incoming water is subject to surface runoff, which is controlled by a user-defined ecoregion parameter (PrecipLossFrac), which is intended to increase with slope or other factors (e.g., rocky soil) that would increase surface runoff even when the soil is not saturated. Incoming precipitation and snowmelt are reduced in proportion to PrecipLossFrac, with the runoff not entering the soil. The actual water infiltrating the soil is:

$$WaterIn = (1 - PrecipLossFrac) \times \frac{[snow\_melt + P - Interception]}{PrecipEvents}$$

### 2.2.2.2 Water Out

Water that infiltrates the soil is subject to both fast and slow “leakage”. Infiltration is limited by the soil saturation parameter ( $\theta_s$ ), and any water in excess of saturation is subject to immediate runoff. Typically, all water above saturation would run off the site, but using the optional RunoffFrac parameter a proportion of the potential runoff can stay on the site creating a supersaturated condition. RunoffFrac values less than 1 will reduce the runoff, and a value of 0 will prevent all runoff.

Fast leakage is correlated to the soil hydraulic conductivity (Ksat) and occurs before plants have a chance to utilize water. The ecoregion parameter LeakageFrac defines the proportion of water above field capacity (-3.37 m pressure head) that immediately drains through the water profile. A parameter value of 1.0 implies immediate draining to field capacity, which will likely be appropriate for most real applications. Values of less than 1 for LeakageFrac could be appropriate to represent soil conditions that prevent leakage through the bottom of the soil profile (e.g., bedrock, clay layer, permafrost). Slow leakage occurs after plant use (transpiration) and evaporation, and keeps the water level at or below field capacity (-3.37 m pressure head) at the end of each monthly time step.

After fast leakage loss has been subtracted, the soil water is subject to further depletion by transpiration and/or evaporation. Transpiration is calculated as the result of plant growth (see Section 2.5). The rate of evaporation is a function of surface radiation (under the canopy), temperature and the soil texture class. Potential evaporation is calculated as a simplified Penmann-Monteith calculation according to Stewart and Rouse (1976) as presented in Cabrera et al. (2016).

Actual evaporation is calculated as:

$$AET = \text{Max}(c \times PET - \text{Transpiration}, 0),$$

where  $c$  is a proportion that decreases linearly from 1.0 when water content is 75% of field capacity, to  $c = 0$  when pressure head is 153 m (i.e., the physical wilting point (Fig 3) (Robock et al. 1995)). AET is limited to the water above the wilting point, so that evaporation ceases when the soil water falls to the wilting point. Transpiration is subtracted from evaporation to reflect decreasing evaporation when the vegetation increases. *De facto* evaporation is 0.0 when LAI > 3.0. Also, surface radiation is automatically 0 when snow cover is present, which results in no evaporation under snow pack.

Transpiration is assumed to use water that otherwise would be subject to evaporation. Therefore, when transpiration exceeds evaporation, no additional water is lost to evaporation.

When some snow melts prior to soil thaw, it runs off and should not become soil water. The best way to account for this (when not modeling soil freezing and permafrost) is to add the proportion of the snowpack lost to runoff to the SnowSublimFrac parameter (section 9.10).

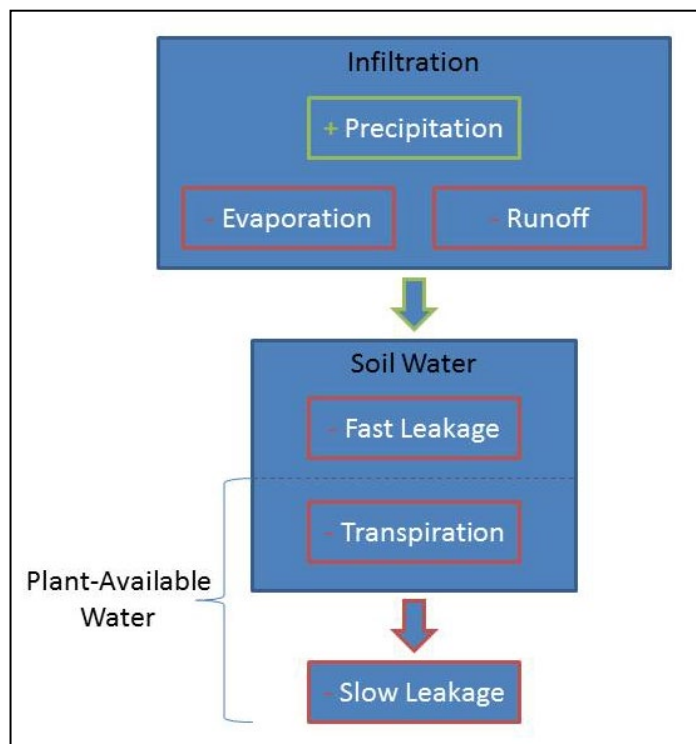


Figure 2. Soil water processes in PnET-Succession

### 2.2.2.3 Water Stress

Water stress in the model depends on the water pressure in the soil according to Feddes et al. (1978). Water pressure in the soil (water retention curves) depends on soil water content and the soil type according to Saxton and Rawls (2006), who provide equations

to estimate water retention curves for soils based on soil texture characteristics (i.e., % sand, % silt, % clay, % organic matter, % gravel, salinity).

Default values for the required parameters (from Saxton and Rawls 2004) are provided with PnET-Succession for 12 different soil types (Figure 3, left panel), but users are able to modify existing soil type parameters or provide custom soil types with parameters. A BEDR (stone) soil type is also included to represent soils with no water holding ability. The user implements a soil type as an ecoregion-specific parameter in the ecoregion parameter table using a corresponding abbreviation for the soil type.

Water stress for a species-cohort is calculated from soil water pressure using four species-specific water pressure thresholds (Figure 3, right panel) labeled H1-H4 in Feddes et al. (1978). Note that PnET-Succession uses the absolute value of pressure head. Parameter H1 (the pressure head below which photosynthesis cannot occur (waterlogging)) is hardcoded in PnET-Succession ( $H1 = 0$  meter pressure). Often, little is known about H2 (cessation of waterlogging stress), so it is recommended to use the generic value  $H2=0.0$  unless you are explicitly modeling waterlogging effects. H3 (onset of stress caused by too little water) can be set to reflect the drought sensitivity of a species, and should fall somewhere between H2 and H4. Most literature sources use a generic H4 (cessation of photosynthesis because of inadequate water) of -153 m pressure head (wilting point).

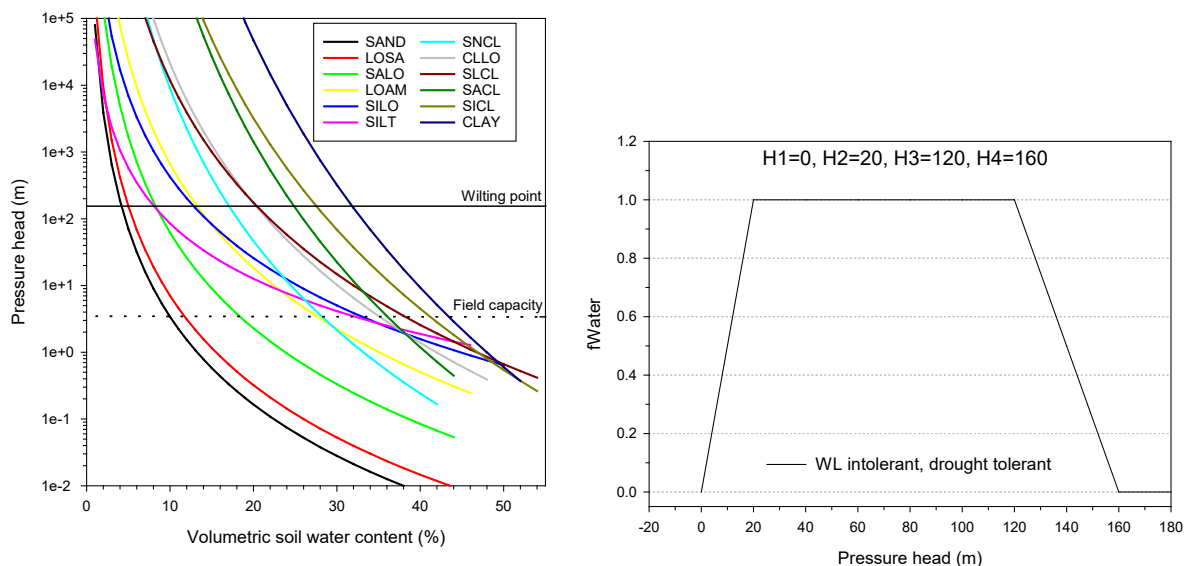


Figure 3. Default pressure head curves (left) and an example of how water stress (fWater, right) is affected by waterlogging and drought tolerance parameters. fWater is calculated by linear interpolation between parameters H1-H4.

Starting with v3.0, water stress (fWater) is used as a reduction factor for the absorption of  $\text{CO}_2$  and  $\text{O}_3$  and the transpiration of water vapor, in addition to its role as a direct growth reduction factor. The internal variable CiModifier is an index of stomatal openness, which is equal to fWater when ozone is not simulated (see 2.4.3.2 for how

ozone affects CiModifier). CiModifier acts as a linear reduction factor in the calculation of the conductance of gases between the atmosphere and the leaves.

## 2.2.3 Other factors

### 2.2.3.1 Temperature

Vapor pressure deficit is calculated from the temperature fluctuations during the day, and accounts for the effect of elevated atmospheric CO<sub>2</sub>. CO<sub>2</sub> affects growth in two ways; 1) it increases water use efficiency and 2) it increases the reference rate of photosynthesis (A<sub>max</sub>). The temperature reduction factor increases from 0 at P<sub>snTMin</sub>, to 1 at P<sub>snTOpt</sub>. Supra-optimal temperatures do not reduce the temperature reduction factor (unless DTEMP=true; see Section 7.19), but net photosynthesis is reduced by VPD effects on conductance and elevated water stress through increased evaporation and transpiration. Foliar respiration is calculated as a user-defined fraction of maximum gross photosynthesis, modified by a temperature reduction factor using a Q-10 relationship. When Wythers=true, foliar respiration is modified to account for acclimation to temperature (see Section 7.18).

### 2.2.3.2 Ozone

Ozone can dramatically reduce photosynthesis rates by damaging photosynthetic tissues and inhibiting stomatal function. The model will simulate these effects when it is optionally supplied with monthly cumulative growing season ozone dose in the climate file. For applications where ozone is not of interest, simply do not supply ozone inputs and no ozone calculations will be made.

The stomatal sluggishness effects of ozone are simulated by altering (increasing) CiModifier (see 2.4.2.3 for water stress influence on CiModifier) to reflect the limited ability of stomata to completely close. Changes to CiModifier (an index of stomatal openness) ripple through the model, altering the CO<sub>2</sub> fertilization effect (DelA<sub>max</sub>), water use efficiency and water stress (through transpiration). Species-specific sensitivity to ozone-induced stomatal sluggishness is controlled by the specification of each species as Sensitive, Intermediate or Tolerant (O<sub>3</sub>StomataSens).

Ozone damage to tissues is simulated using an ozone growth reduction factor (fO<sub>3</sub>) that reduces A<sub>max</sub> to reflect decreased photosynthetic capacity. Species-specific sensitivity to ozone-induced growth reduction is controlled by a scaling parameter (O<sub>3</sub>GrowthSens) that scales the ozone effect to be higher or lower than the average effect. Technical details of the new ozone capabilities can be found in Gustafson et al (in review). Contact Eric Gustafson ([egustafson@fs.fed.us](mailto:egustafson@fs.fed.us)).

## 2.3 Cohort Growth and Ageing

The model computes net photosynthesis, and gross photosynthesis is estimated by adding foliar respiration to net photosynthesis. Gross photosynthesis is used to compute transpiration. Net photosynthesis production is allocated to maintenance respiration and then to the root, foliage, wood and non-structural biomass pools, according to fixed allocation ratios. A proportion of foliage and wood biomass is also



moved to the dead pools to simulate leaf-fall and stem (self-thinning)/branch/root death. Cohort ageing is simply the addition of the time step to each existing cohort age.

Cohort wood biomass and foliage can be directly altered by disturbance extensions. Partial disturbances can remove portions of total biomass. Defoliation disturbances can directly remove foliage, without altering the wood biomass. The loss of foliage has subsequent consequences for photosynthetic capacity due to less leaf area. Starting with v3.4, cohorts can re-flush foliage following defoliation. Refoliation occurs at 70% of the calculated ideal foliage, for deciduous cohorts ( $TO_{fol} = 1$ ) that experience  $> 60\%$  defoliation in a given year. Refoliation has additional cost to NSC (95% of ideal foliage), and cohorts that do not reflate still experience additional NSC losses (10% of ideal foliage).

## 2.4 Cohort Senescence and Mortality

Senescence is implemented as a reduction of gross photosynthetic rate with age such that respiration eventually exceeds production and cohorts die. A cohort dies when non-structural carbon decreases to  $< 1\%$  of the combined structural biomass pools. The  $PsnAgeRed$  parameter controls the shape of the function used to calculate the age-related reduction factor, which reaches zero at the longevity specified in the LANDIS-II species parameter file. Cohorts are also removed and considered dead when their biomass falls below 1 g/m<sup>2</sup>.

## 2.5 Dead Biomass Decay

When a cohort dies and is removed (e.g., harvest or insect mortality), its biomass is added to one or both of the dead biomass pools: woody and leaf. Decomposition rate of woody litter depends on a decay rate that is weighted by additions of woody material and user-supplied species specific decay rates ( $KWdLit$ ). Decomposition rate of non-woody litter depends on a weighted decay rate according to additions of foliage and their associated decomposition rates that depend on species specific foliage lignin concentrations ( $FolLignin$ ) and ecosystem determined AET according to Meentemeyer (1978). Disturbances can alter the dead biomass pools. They can add dead biomass (e.g., wind) and/or remove dead biomass (e.g., fire may add some woody dead biomass and remove all leaf dead biomass).

## 2.6 Cohort Reproduction and Establishment

Cohort establishment is the result of two distinct processes: 1) production and dispersal of seeds and 2) seed germination and successful recruitment of a viable new cohort.

Seed is produced by every cohort that is at least the age of sexual maturity. Seed dispersal is modeled as a spatial process according to the dispersal method selected by the user, as in the Biomass Succession extension.

When seeds disperse to a cell, establishment (recruitment) is stochastic based on a probability of establishment that is calculated as a function of soil moisture and sub-canopy radiation during the time step. Establishment is only attempted once per timestep, using the average light and water conditions during optimal months, up to three physiologically active months in the year, based on temperature. The first active

month each year is not used for establishment calculations, assuming trees put resources into leaf and stem growth in the first month, and seeds first become available in the second month.

Establishment probability for a give timestep is generally calculated with the following equations:

$$fRad = \frac{SubcanopyPAR}{SubcanopyPAR + HalfSat}$$

$$fRadAdj = fRad^2 \times \frac{1}{EstRad^2}$$

$$fRadAdjInt = \frac{HalfSat - MinHalfSat}{HalfSatRange}$$

$$fRadAdjSlope = (fRadAdjInt \times 2) - 1$$

$$fRadFinal = 1 - fRadAdjInt + fRadAdj \times fRadAdjSlope$$

$$fWaterAdj = fWater^2 \times \frac{1}{EstMoist^2}$$

$$p_{Est} = fWaterAdj \times fRadFinal \times MaxPest$$

Where MinHalfSat is the lowest HalfSat among all modeled species, HalfSatRange is the range of species HalfSat values, and fWater is calculated from available water as described in 2.2.2.3. The establishment modifiers (EstRad and EstMoist) are used to scale the relative impacts of light and water stress on the probability of cohort establishment. EstRad specifies the fRad above which the light environment is optimal for establishment, and EstWater specifies the fWater above which the soil moisture is optimal for establishment.

Initial biomass for a successfully established cohort is computed for a 1-year old cohort. This initial cohort will be grouped ('binned') appropriately into a larger cohort (e.g., age 1 – 10) at the next succession time step.

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### 3 Input File - PnET-Succession

The input parameters for this extension are specified in two primary input files: the PnET-Succession input file and the PnET Species Parameters input file. The general species parameter input file used by all versions of LANDIS is also required, and is described in Chapter 6 of the *LANDIS-II Model User Guide*. The input files must comply with the general format requirements described in section 3.1 *Text Input Files* in the *LANDIS-II Model User Guide*.

#### 3.1 Example PnET-Succession input file

```
LandisData  "PnET-Succession"

PnET-Succession      Value
>>-----
Timestep              10
StartYear             1961
SeedingAlgorithm      WardSeedDispersal
MaxDevLyrAv           6000

PNEToutputsites       PNETOutput.txt

InitialCommunities    Oconto_initial-communities.txt
InitialCommunitiesMap Oconto_initial-communities.img

PnETGenericParameters PnETGenericParameters.txt
PnETSpeciesParameters PnET_Oconto_species.txt
EcoregionParameters   Oconto_EcoregionParameters.txt
DisturbanceReductions  disturbance_reductions.txt
ClimateConfigFile      climate-generator.txt
```

#### 3.2 LandisData

This parameter's value must be "PnET-Succession".

#### 3.3 Timestep

This parameter is the time step of the extension. A value <5 is recommended. Random shuffling of cohort foliage into sub-canopy layers for access to light is done at each time step, so a poor random assignment in a long time step may kill cohorts that would survive with a shorter time step. Longer time steps do not markedly speed up simulations or reduce output because the internal time step of PnET-Succession is monthly, but they do reduce the frequency of outputs. Value: integer >0. Units: years.

#### 3.4 StartYear

This parameter indicates the climate year in which simulation begins. Climate file observations prior to this date are used for spin-up (as necessary) and observations from this date forward are used for simulations. The climate file may contain more years than will actually be used by the model. Value: integer > 0. Units: years.

### 3.5 SeedingAlgorithm

This parameter is the seed dispersal algorithm to be used. Valid values are "WardSeedDispersal", "NoDispersal" or "UniversalDispersal". The algorithms are described in section 4.5.1 *Seeding of the LANDIS-II Conceptual Model Description*.

### 3.6 PNETOutputsites

Optional: Invoke the output extension PNETOutputsites and specify its input file (see section 10).

### 3.7 InitialCommunities

This parameter gives the name of the initial communities text file. This file assigns species and cohorts to each value found in the initial communities map (see section 4).

### 3.8 InitialCommunitiesMap

This parameter gives the file name of the initial communities map. This map contains a unique integer value for each combination of species and cohorts found on the landscape. Each cell value for an active site on the landscape must be one of the map codes listed in the initial communities input file (see section 5).

### 3.9 LitterMap (Optional)

This parameter gives the file name of the initial leaf litter map. The values of the map represent the quantity ( $\text{g/m}^2$ ) of leaf litter on each site at the start of the simulation. When providing initial cohort biomass values, it is valuable to also provide initial litter values, which will otherwise start with 0 values. When initial cohort biomass is not provided (estimated from spin-up), then the litter quantity is also estimated during the spin-up phase.

### 3.10 WoodyDebrisMap (Optional)

This parameter gives the file name of the initial dead woody debris map. The values of the map represent the quantity ( $\text{g/m}^2$ ) of dead wood on each site at the start of the simulation. When providing initial cohort biomass values, it is valuable to also provide initial dead wood values, which will otherwise start with 0 values. When initial cohort biomass is not provided (estimated from spin-up), then the dead wood quantity is also estimated during the spin-up phase.

### 3.11 PnETGenericParameters

This optional parameter gives the name of a PnET Generic Parameter text file. Any parameter that is not species-specific, or is typically specified in the PnETSpeciesParameter file (Chapter 8), but is identical for all species can be supplied either in the default generic parameter file installed in C:\Program Files\LANDIS-II\v6\bin\extensions\Defaults\GenericPnETSpeciesParameters.txt with the rest of the model, or in a custom generic parameter file specified here. Any parameters not specified in the PnETSpeciesParameter file will be read from the custom generic file,

and if not found there, will be read from the default generic file. Thus, values found in the PnETSpeciesParameter file will always take precedence over the default generic file, but cannot be duplicated in the PnETGenericParameters file. The format of the PnET Generic Parameter text file is described in section 7.

### 3.12PnETSpeciesParameters

This parameter gives the name of the PnET Species Parameter text file. The format of this file is described in section 8.

### 3.13EcoregionParameters

This parameter gives the name of the PnET Ecoregions Parameter text file, which is described in section 9.

### 3.14DisturbanceReductions (Optional)

This parameter gives the name of the Disturbance Reductions txt file, which is described in Section 10. If this parameter is not supplied, all disturbances will cause 100% of cohort biomass to contribute to the dead pools, and dead pools will not be reduced by any disturbances.

### 3.15ClimateConfigFile (Optional)

This optional parameter gives the name of the Climate Library configuration file. Including this optional parameter triggers the model to use the climate library to provide the climate variables used within PnET-Succession. Excluding this parameter will utilize the climate input file(s) referenced in the EcoregionParameters input file, which is described in Section 9.13.

### 3.16SaxtonAndRawlsParameters (Optional)

This optional parameter gives the name of the soil parameter file should the user wish to use a file other than the default provided installed with PnET-Succession.

### 3.17CohortBinSize (Optional)

This optional parameter sets the number of years represented by an age cohort. Typically, the age range of a cohort is determined by the timestep of the succession extension. However, in cases where users want to lump the cohorts into combined age cohorts, while maintaining a more frequent succession timestep, this parameter can be used. The CohortBinSize must be  $\geq$  Timestep.

## 4 Input File – Initial community classes

This file contains the definitions of the initial community classes. Each active site on the landscape is assigned to an initial community class. The class specifies the tree species that are present along with the particular age cohorts that are present for each species. Avoiding a proliferation of similar-aged cohorts of the same species on a site will speed run times with little effect on simulation results because those cohorts would be competing with each other.

Note: The format of the text file produced by the Biomass Community Output extension is compatible to be used as an input initial community file.

### 4.1 Example File

```
LandisData "Initial Communities"

>>Old jackpine oak
MapCode 7
    acerrubr 30
    pinubank 90
    pinuresi 110 140
    querelli 40 240

>> young jackpine oak
MapCode 0
    pinubank 50
    querelli 10 70

>> young aspen
MapCode 2
    poputrem 10

>> old maple hardwoods
MapCode 55
    abiebals 10 60 120
    acerrubr 90 120
    acersacc 20 50 150 200
    betualle 40 140 200
    fraxamer 10 100 180
    piceglau 180
    querrubr 100 160 180
    thujocci 200 260
    tiliamer 20 80 150
    tsugcana 30 80 120 220 320 340

>> old pine - spruce - fir
MapCode 6
    abiebals 10 50 80
    piceglau 100 180 220
    pinuresi 140 180
    pinustro 200 280 350
```

### 4.2 LandisData

This parameter's value must be "Initial Communities".

## 4.3 Initial Community Class Definitions

Each class has an associated map code and a list of species present at sites in the class.

### 4.3.1 MapCode

This parameter is the code used for the class in the input map (see chapter 5). Value:  $0 \leq \text{integer} \leq 65,535$ . Each class map code must be unique. Map codes can appear in any order, and need not be consecutive.

### 4.3.2 Species Present

A list of species present at the class' sites comes after the map code. Each species is listed on a separate data line.

```
species age age age ...
```

The species name comes first, followed by one or more ages. The name and ages are separated by whitespace. An age is an integer and must be between 1 and the species' Longevity parameter. The ages can appear in any order.

```
acersacc 10 5 21 60 100
```

The list may be empty, which will result in the sites in the class being initialized with no species cohorts.

#### 4.3.2.1 Cohort Biomass (optional)

Initial community information can also provide initial biomass values for cohorts. Supplying initial biomass values is optional, and when not provided the extension will use the succession grow functions to grow the cohort to the initial age. Biomass values should represent aboveground live woody biomass ( $\text{g/m}^2$ ), and should be provided in parentheses following the age

```
acersacc 10 (2100) 60 (5200) 100 (8500)
```

Biomass values should be greater than 0. A cohort assigned a biomass of 0 will be treated as if no biomass value was provided, and will have its biomass estimated through the spin-up process.

### 4.3.3 Grouping Species Ages into Cohorts

The list of ages for each species is grouped into cohorts based on the succession extension's timestep or the optional parameter CohortBinSize. This value determines the size of each cohort. For example, if the timestep is 20, then the cohorts are ages 1 to 20, 21 to 40, 41 to 60, etc.

Suppose an initial community class has this species in its list:

```
acersacc 10 25 30 40 183 200
```

If the succession timestep is 10, then the cohorts for this species initially at each site in this class will be:

```
acersacc 10 20 30 40 190 200
```



## 5 Input File – Initial community map

This is a GIS file of the initial community classes. Each active site on the landscape is assigned to a MapCode that links to the initial community class defined in the Initial Community Class Definitions. The file can be in any valid LANDIS-II map format.

Note: The format of the map file produced by the Biomass Community Output extension is compatible to be used as an input initial community map file.

## 6 Input File – Climate

This file contains weather records of monthly parameter values. Note that PnET-Succession can optionally access the LANDIS-II climate library although a climate file is still required to prevent an error.

### 6.1.1 Example File #1

Year	Month	Tmax	Tmin	Prec	PAR	CO2
1700-1979	1	1.57	-7.86	96.59	493.10	335
1700-1979	2	3.46	-6.94	87.36	671.21	335
1700-1979	3	8.54	-3.16	110.79	852.52	335
1700-1979	4	15.50	2.34	110.38	925.71	335
1700-1979	5	20.37	7.69	133.10	873.77	335
1700-1979	6	24.39	12.38	123.72	872.04	335
1700-1979	7	26.28	14.81	135.81	847.05	335
1700-1979	8	25.64	14.00	109.89	842.81	335
1700-1979	9	22.07	10.13	100.57	760.33	335
1700-1979	10	16.06	4.03	89.03	624.86	335
1700-1979	11	9.83	-0.88	101.25	463.20	335
1700-1979	12	3.77	-5.33	100.48	411.67	335
1980	1	1.57	-7.21	53.32	496.07	338
1980	2	-0.37	-9.99	47.78	697.67	338
1980	3	6.85	-4.81	133.22	857.97	338
1980	4	14.52	2.06	139.30	907.75	338
1980	5	21.07	7.89	122.41	927.89	338
1980	6	23.51	9.96	137.54	925.58	338
1980	7	26.88	15.04	154.32	818.81	338
1980	8	26.42	15.58	169.70	799.45	338
1980	9	23.97	11.23	65.91	797.72	338
1980	10	14.23	2.19	76.10	634.71	338
1980	11	7.82	-2.33	93.34	445.37	338
1980	12	3.12	-6.90	47.77	468.97	338

### 6.1.2 Example File #2

Year	Month	Tmax	Tmin	PAR	Prec	CO2	O3
1995	1	-4.22	-13.70	262.4	9.9	367	0.0
1995	2	-4.21	-14.7	487.9	13	367	0.0
1995	3	5.08	-6.67	1012.02	35.9	367	0.0
1995	4	7.58	-3.68	741.21	49.3	367	0.0
1995	5	18.37	5.12	1021.65	103.4	367.7	2769
1995	6	27.26	12.67	1292.2	21.5	362.8	3279
1995	7	26.37	14.15	1287.3	137.2	367.9	3589
1995	8	25.87	15.41	1118.77	145.8	372.5	5686
1995	9	19.55	6.24	770.17	65.6	364.9	7723
1995	10	12.10	2.72	531.73	138	368	8167
1995	11	-1.58	-10.74	299.5	51.6	368	0.0
1995	12	-5.45	-15.41	233.53	34.5	368	0.0

## 6.2 Header Information

The first line of the file must contain the following text:

```
Year   Month   TMax  Tmin   PAR    Prec   CO2    O3 <optional>
```

## 6.3 Observations

Subsequent lines of the file contain monthly values for each variable. Observations must appear in chronological order.

### 6.3.1 Year

The year of the weather observation. Alternatively, a range of years may appear, delineated by a hyphen (see example 6.1.2). Value: 4-digit integer >0.

### 6.3.2 Month

The month of the weather observation. Value:  $1 \leq \text{integer} \leq 12$ .

### 6.3.3 TMax

The maximum temperature observed in the month. Value: decimal. Units: degrees C.

### 6.3.4 Tmin

The minimum temperature observed in the month. Value: decimal. Units: degrees C.

### 6.3.5 PAR

Mean monthly value of Photosynthetically Active Radiation during daylight hours. Value: decimal  $\geq 0.0$ . Units: User choice. Typically  $\mu\text{mol}/\text{m}^2/\text{sec}$  or  $\text{W}/\text{m}^2$ . The

units for the half-saturation constant (SpeciesParameter file) must be the same as PAR. **THE MODEL WILL NOT CHECK TO ENSURE THAT THE UNITS ARE THE SAME.** This is a user responsibility.

### 6.3.6 Prec

The sum of precipitation observed in the month. Value: decimal  $\geq 0$ . Units: mm.

### 6.3.7 CO2

Mean monthly atmospheric CO<sub>2</sub> concentration. Value: decimal  $> 0$ . Units: ppm.

### 6.3.8 O3 (Optional)

Cumulative atmospheric ozone (O<sub>3</sub>) dose over 40 ppb (CumD40) for the growing season. Computed by subtracting 40 from each hourly O<sub>3</sub> value and summing non-negative values for all hours between 0800 and 1900 hours in the month. Each monthly total is added to the previous monthly total to give the cumulative growing season hourly dose through the current month. Value: decimal  $\geq 0$ . Units: ppb-h.

## 7 Input File – Generic PnET Species Parameters

This file contains PnET parameters that are not species-specific or are identical for all species. Only parameters that are not described in Chapter 8 are described here. Parameters may appear in any order. **NOTE:** Any of these parameters may instead be set in the PnETGenericDefaultParameters.txt file in the Defaults folder found where PnET-Succession is installed (usually in C:\Program Files\...\extensions\). Parameters set here will over-ride settings in the PnETGenericDefaultParameters file. Also, most (but not all) of these parameters may instead appear in the PnETSpeciesParameter file (Section 8), but may not appear in both files.

### 7.1 Example file:

```
LandisData PnETGenericParameters

PnETGenericParameters Value
>>-----
MaxCanopyLayers      2
MaxDevLyrAv          6000
IMAX                 10
PreventEstablishment true
DVPD1                0.05
DVPD2                2
BFolResp             0.1
MaintResp            0.002
TOroot               0.02
TOwood               0.01
Q10                  2
FolLignin            0.2
KWdLit               0.01
InitialNSC            7
CFracBiomass         0.45
```

PrecipEvents	11
Wythers	true
DTEMP	true

## 7.2 LandisData

This parameter's value must be "PnETGenericParameters".

## 7.3 PnETGenericParameters

This keyword must be followed by "Value".

## 7.4 MaxCanopyLayers

Optional parameter that caps the number of canopy layers that can be implemented. Typically, forest canopy layers will not exceed 5 and applying many canopy layers slows the model dramatically. The number of canopy layers should primarily be regulated through MaxDevLyrAv, but when MaxDevLyrAv is given low values, the result would otherwise be an extreme number of canopy layers. MaxCanopyLayers has a default value of 5.

## 7.5 MaxDevLyrAv

This optional parameter is used to lump species-age cohorts into canopy layers, and specifies the maximum variation of cohort biomass that can occur within a canopy layer. It is given a large default value, which results in a single canopy layer regardless of biomass distribution among subcanopy layers (see section 2.4.1).

## 7.6 IMAX

Optional: Each cohort is subdivided into a number of layers (IMAX) for integration. In PnET (Aber and Federer, 1992), the number of subcanopy layers was fixed at 50. Reducing IMAX saves computation time, with robust results when  $IMAX \geq \sim 5$  (De Bruijn et al. 2014). When omitted, the model uses the default  $IMAX=10$ .

## 7.7 DVPD1, DVPD2

Coefficients for converting vapor pressure deficit (VPD) to DVPD according to  $DVPD = 1 - DVPD1 * vpd^{DVPD2}$  (photosynthesis reduction factor due to vapor pressure). Value: decimal. Units: kPa<sup>-1</sup>.

## 7.8 BFolResp

Base Foliar Respiration Fraction - Foliar respiration as a fraction of maximum photosynthetic rate. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion.

## 7.9 MaintResp

Loss of NSC due to maintenance respiration. This rate is modified by temperature using a Q10 relationship, and is applied to the biomass pool monthly. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion of NSC lost per month.

### 7.10TORoot/TOWood

Turnover of Root/Wood - Fraction of root/wood biomass lost per year to damage, breakage or death. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion per year.

### 7.11Q10

Respiration  $Q_{10}$  value for foliar respiration, a measure of the rate of change of respiration when temperature is increased by 10 °C. Value:  $0.0 \leq \text{decimal} \leq 10.0$ . Units: none.

### 7.12FolLignin

Mass fraction of lignin in foliage tissue. Value:  $0.0 \leq \text{decimal} \leq 0.8$ . Units: gr/gr.

### 7.13KWdLit

Annual decomposition rate (decay constant,  $k$ ) of woody litter. Value:  $0.0 \leq \text{decimal} \leq 0.4$ . Units: proportion per year.

### 7.14InitialNSC

Amount of NSC assigned to newly established cohorts. Value: integer > 0. Units: g.

### 7.15CFracBiomass

Carbon fraction of biomass by weight. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion.

### 7.16PrecipEvents

Monthly total precipitation is evenly divided among this number of events to allow cohorts more opportunities to compete stochastically for incoming water. Lower values tend to reduce the competitiveness of species with lower H3/H4 values. Default=11. Value: decimal  $\geq 1.0$ . Units: count.

### 7.17PreventEstablishment

Boolean variable turning establishment on or off. Value= “true” or “false”.

### 7.18Wythers

Boolean variable turning the Wythers correction on or off. The Wythers algorithm accounts for acclimation of foliar respiration to elevated temperatures (Wythers et al 2013). Value: true/false. Default: false.

### 7.19DTEMP

Boolean variable turning the PnET-II DTEMP temperature reduction factor (Aber and Federer 1992) on or off. A value of “true” uses the DTEMP function, which reduces NetPsn from 1.0 at PsnTOpt to 0.0 at both PsnTMin and PsnTMax. A value of “false”

will use the original PnET-Succession (v1.0) function that behaves the same as DTEMP at temperatures below PsnTOpt, but does not reduce NetPsn at temperatures above PsnTOpt, relying solely on VPD-driven reductions to drop NetPsn to zero at about 40 °C. DTEMP produces somewhat more of a NetPsn penalty above PsnTOpt for species with PsnTOpt <~24 °C. The difference between the functions is small for species with PsnTOpt >24 °C. Value: true/false. Default: false.

## 7.20 InvertPest (optional)

Boolean variable activating the optional inversion of the effect of fRad on Pest as a function of HalfSat (see section 1.3.1). Values: true or false. Default: false.

## 7.21 Permafrost (optional)

Boolean variable activating the optional simulation of soil ice depth and its effects on leakage. Not recommended unless permafrost exists in your study area because it slows runtimes. Values: true or false. Default: false.

## 7.22 LeakageFrostDepth (optional)

Soil ice depth at which frozen soil no longer impedes soil drainage. LeakageFrac gradually increases from 0.0 when ice depth is ≤RootingDepth to 1.0 at LeakageFrostDepth. Can be omitted when Permafrost=false. Value: integer>0. Units: mm.

# 8 Input File – PnET Species Parameters

These parameter values typically vary by species. If they do not, they may more conveniently be placed in the PnETGenericParameters file (Section 7). All parameters for a species appear on a single line. Parameters may appear in any order.

## 8.1 Example file:

LandisData PnETSpeciesParameters

PnETSpeciesParameters	FolN	SLWmax	SLWDel	TOfol	AmaxA	AmaxB	HalfSat	H3	H4	PsnAgeRed			
PsnTMin PsnTOpt k	DNsc	FracBelowG		EstMoist		EstRad	FracFol	FrActWd	CO2HalfSatEff	O3StomataSens			
O3GrowthSens	FolNInt	FolNSlope											
abiebal 1.5	160	0	0.25	5.3	21.5	150	150	275	5	2	19	0.5	0.05
0.25 10	3	0.053	0.00002	0.5	Sensitive	2.0	0.8	0.4					
acerrub 2.5	65	0.2	1	-46	71.9	200	150	500	5	4	26	0.58	0.05
0.33 5	5	0.028	0.00004	0.0	Sensitive	2.0	0.5	1.0					
acersac 2.4	50	0.2	1	-46	71.9	100	150	275	5	2	23	0.58	0.05
0.33 10	1	0.02	0.00002	0.0	Tolerant	0.0	0.8	0.4					

## 8.2 LandisData

This parameter's value must be "PnETSpeciesParameters".

### 8.3 PnETSpeciesParameters (species name)

The species name as it appears in the species parameter input file (see Chapter 6 of the *LANDIS-II Model User Guide*).

### 8.4 FolN

Foliar nitrogen content (%). Value:  $0 < \text{decimal} < 10$ . Units: %.

### 8.5 SLWmax

Maximum specific leaf weight at the top of canopy. Value:  $0 < \text{decimal} < 1000$ . Units:  $\text{g/m}^2$ .

### 8.6 SLWDel

Rate of change in specific leaf weight from the top of a canopy layer to the bottom. Set to zero to make SLW constant throughout a canopy layer. Value:  $0.0 \leq \text{decimal} \leq 2$ . Units:  $\text{g}^{-1}\text{fol}$ .

### 8.7 Tofol

Turnover of foliage - Fraction of foliage biomass lost per year. Typically the reciprocal of leaf longevity. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion per year.

### 8.8 AmaxA

Intercept of relationship between foliar N and maximum net photosynthetic rate. Units:  $\text{nmol CO}_2 \text{ g}^{-1} \text{ leaf s}^{-1}$ . Value:  $-500 < \text{decimal} < +500$

### 8.9 AmaxB

Slope of relationship between foliar N and maximum net photosynthetic rate, such that  $A_{\text{max}} (\text{nmol CO}_2 \text{ g}^{-1} \text{ leaf s}^{-1}) = A_{\text{maxA}} + A_{\text{maxB}} * \text{FoliarN}$ . Units  $\text{nmol CO}_2 \text{ g}^{-1} \text{ leaf s}^{-1}$ . Value: decimal  $> 0$ .

### 8.10 HalfSat

Half saturation light level for photosynthesis. Lower values reflect more shade tolerance. Value: integer  $> 0$ . Units: User choice. Typically  $\mu\text{mol/m}^2/\text{sec}$  or  $\text{W/m}^2$ . The units of PAR in the climate input file must be the same as HalfSat. **THE MODEL WILL NOT CHECK TO ENSURE THAT THE UNITS ARE THE SAME.** This is a user responsibility. Note: HalfSat is the only parameter that determines shade tolerance in PnET-Succession; the ShadeTolerance parameter required in the LANDIS-II species file is ignored by PnET-Succession.

### 8.11 H1, H2, H3, H4

Water stress parameters according to Feddes et al. (1978). See Section 2.4.2.3. H1 may take a negative value to produce  $f_{\text{Water}} > 0.0$  when pressure head = 0 (waterlogging tolerance). H1, H2, H3 and H4 should be successively larger positive

values. Note that this is the absolute value of actual pressure head values. Units: m pressure head.

### 8.12PsnAgeRed

Reduction factor reducing leaf photosynthesis rate as cohorts age, with fAge=1.0 at age 1 and fAge =0.0 at the longevity specified in the LANDIS-II species parameter file. Longevity should be specified as longevity under optimal conditions because the various reduction factors will combine to almost always result in cohort death prior to the specified longevity age. A value <1.0 results in a rapid initial decline in max photosynthesis with age, a value of 1.0 results in a linear decline and a value >1.0 results in slow initial decline, according to:  $y = \left( \frac{age}{longevity} \right)^{PsnAgeRed}$ . Cohorts die when NSC is <1% of the value of the other biomass pools combined at the end of a calendar year. Value: 0.0< decimal <infinity. Units: proportion per year.

### 8.13LeafOnMinT (Optional)

Minimum temperature for active growing season. Compared to Tmin to determine start and end month of growing season. Foliage is allocated during all active months. If this parameter is not provided, the value of PsnTMin is used in its place (same as previous version). Value: decimal. Units: °C.

### 8.14PsnTMin

Minimum average daytime temperature for photosynthesis. Value: decimal. Units: °C.

### 8.15PsnTOpt

Optimal average daytime temperature for photosynthesis. Value: decimal >PsnTMin. Units: °C.

### 8.16PsnTMax (optional)

Maximum average daytime temperature for photosynthesis. Typically not greater than 37°C. Value: decimal >PsnTOpt. Units: °C.

If PsnTMax is not provided, an estimated value of PsnTMax is calculated as (PsnTOpt + PsnTOpt – PsnTMin), which is consistent with earlier versions of the model.

### 8.17ColdTol (optional)

Cold tolerance. Coldest temperature that the species can survive. Value: decimal. Units: °C.

### 8.18k

Canopy light attenuation constant (light extinction coefficient). Value:  $0.0 \leq$  decimal  $\leq 1.0$ . Units: none.



## 8.19DNSC

Proportion of NSC relative to total active biomass that will be maintained as long as net photosynthesis exceeds maintenance respiration. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion of active biomass.

## 8.20FracBelowG

Fraction of non-foliar biomass that is belowground (root pool). Allocations vary at each time step to maintain this fraction. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion.

## 8.21EstMoist

Scaling parameter that causes the maximum expected value of fWater to equal 1.0 when computing Pest. Because fWater usually equals 1.0 at some time in a season or simulation, this parameter is typically set equal to 1.0. Purpose is to make light and water have an equal effect on establishment. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion.

## 8.22EstRad

Scaling parameter that causes the maximum expected value of fRad to equal 1.0 when computing Pest. Because fRad almost never equals 1.0 during a simulation, this parameter is typically set equal to the maximum fRad value observed in the first year when a single cohort of a species is grown on an open site. Purpose is to make light and water have an equal effect on establishment. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion.

## 8.23MaxPest

Tuning parameter used to raise or lower the Pest of all species by the same proportion. EstMoist and EstRad equalize the effect of water and light, linking these effects to the tolerance of the species to each factor, and MaxPest is used to control overall cohort proliferation. Note that this parameter gives the maximum Pest for the time step; for different values of Timestep, MaxPest should vary in proportion to Timestep. It is also affected by CohortBinSize, so MaxPest should be calibrated for a specific combination of Timestep and CohortBinSize. It is not unusual for this parameter to take values  $<0.5$  when Timestep  $<10$ , and it may be  $<0.1$  when Timestep = 1. Value:  $0.0 \leq \text{decimal}$ . Units: proportion.

## 8.24FracFol

Fraction of the amount of active woody biomass (above and belowground) that is allocated to foliage each year. The active fraction of woody biomass is calculated by the model using FracActWd. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion per year.

## 8.25FrActWd

Shape parameter of negative exponential function that calculates the amount of woody biomass that has active xylem capable of supporting foliage. All wood is active when

the parameter =0.0, and increasing values decrease the fraction of active wood as biomass increases according to:  $\text{active\_wood} = e^{-(\text{FrActWd} * \text{biomass})}$ . Value:  $0.0 \leq \text{decimal} \leq 0.4$ . Units: unitless.

### 8.26CO2HalfSatEff (Optional)

Slope coefficient used to reduce HalfSat (increase shade tolerance) as CO<sub>2</sub> concentration increases. Value:  $\text{decimal} \leq 0.0$ . Units: unitless. Default=0.0. Set to zero to turn off this effect. See PnET-Succession function worksheet.xlsx to see how the parameter affects HalfSat.

### 8.27O3StomataSens (Optional)

Categorical parameter indicating the species' susceptibility of the stomata to ozone-induced sluggishness. Values: one of – Sensitive, Intermediate, Tolerant.

### 8.28O3GrowthSens (Optional)

Scaling parameter controlling the species' susceptibility to ozone-induced tissue damage. Zero indicates no sensitivity to ozone, 2.0 indicates twice as much sensitivity as 1.0, 3.0 indicates three times as much sensitivity as 1.0. Value:  $0.0 \leq \text{decimal}$ . Units: unitless. Default=0.0. Set to zero to turn off effect.

### 8.29MaxFolN, FolNShape (Optional)

Parameters controlling the dynamic response of foliar nitrogen (FolN) to light according to:

$$\text{AdjFolN} = \text{FolN} + ((\text{MaxFolN} - \text{FolN}) * (\text{fRad}^{\text{FolNShape}}))$$

FolN is used as the minimum bound for AdjFolN. Values:  $0.0 \leq \text{decimal}$ . Units: %N by weight. Omit these parameters to turn off effect. See PnET-Succession function worksheet.xlsx to see how the parameters affect FolN.

### 8.30MaxFracFol, FracFolShape (Optional)

Parameters controlling the dynamic response of fraction of foliage (FracFol) to light according to:

$$\text{AdjFracFol} = \text{FracFol} + ((\text{MaxFracFol} - \text{FracFol}) * (\text{fRad}^{\text{FracFolShape}}))$$

FracFol is used as the minimum bound for AdjFracFol. Values:  $\text{decimal} \geq 0.0$ . Units: proportion, unitless. Defaults=1.0, 0.0. Omit these parameters to turn off effect. See PnET-Succession function worksheet.xlsx to see how the parameters affect FracFol.

## 9 Input file - Ecoregion parameters

### 9.1 Example file:

LandisData      EcoregionParameters

EcoregionParameters SoilType Latitude RootingDepth PrecLossFrac LeakageFrac PrecIntConst SnowSublimFrac  
ClimateFileName

eco1 SALO 46 1000 0.0 1.0 0.11 0.15 Climate-input.txt

### 9.2 LandisData

This parameter's value must be "EcoregionParameters".

### 9.3 EcoregionParameters (ecoregion name)

The ecoregion name given must be defined in the ecoregion input file (see chapter 7 in the *LANDIS-II Model User Guide*). Ecoregions may appear in any order.

### 9.4 SoilType

Abbreviation for the predominant soil type in the ecoregion. Soil type is used in the model to determine the water retention curve of the soil (see Figure 3 and section 2.4.2.3). Soil type names can be any character string that also appears in the corresponding soil parameters file (SaxtonAndRawlsParameters). A default version of this parameter file installs with the extension (C:\Program Files\LANDIS-II\v6\bin\extensions\Defaults), and supports the following soil types: SAND (sand), LOSA (loamy sand), SALO (sandy loam), LOAM (loam), SILO (silt loam), SILT (silt), SNCL (sandy clay loam), CLLO (clay loam), SLCL (silty clay loam), SACL (sandy clay), SICL (silty clay), CLAY (clay), BEDR (stone). These categories correspond with FAO soil types. Value: 4-letter string, case sensitive.

### 9.5 Latitude

This parameter is the approximate latitude of the ecoregion, used to compute daylength. Value:  $-90 < \text{integer} < 90$ . Units: degrees of latitude.

### 9.6 RootingDepth

This represents the rooting zone, the soil depth to which roots typically penetrate. Using the bucket-model analogy, this determines the depth of the "bucket." Value:  $\text{integer} \geq 0$ . Units: mm.

### 9.7 PrecLossFrac

Precipitation Loss Fraction. Proportion of precipitation that does not enter the soil (e.g., runoff induced by topographic slope). Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion.

## 9.8 LeakageFrac

Leakage Fraction. Proportion of soil water above field capacity that is subject to “fast leakage” (see Figure 2). Fast leakage is the drainage of infiltrated water that leaks out of the rooting zone soil more quickly than plants can access the water. Therefore, water lost to fast leakage is not available for transpiration. A value of 1.0 for LeakageFrac will cap plant available water at field capacity. Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion.

## 9.9 RunoffCapture

Height above soil surface that water can reach before any runoff occurs, representing height of basin outlet. This parameter is used to allow soil water to accumulate above saturation (i.e., flooded). RunoffCapture only applies to water that exceeds the soil porosity (saturation), and would cause it to all run off when RunoffCapture = 0 (the previous model behavior). Any excess water (above saturation) is retained in the soil “bucket” until evaporation, interception, transpiration and leakage (if any) collectively exceeded cumulative precipitation inputs (i.e., consumed all excess water). Value:  $\text{decimal} \geq 0.0$ . Units: mm.

## 9.10 PrecIntConst

This represents the rate of precipitation interception for each unit of leaf area index, which is lost to evaporation, and therefore does not enter the soil. See the interception equation in section 2.4.2.1. Value:  $\text{decimal} \geq 0.0$ . Units: unitless.

## 9.11 SnowSublimFrac

Fraction of the snowpack that sublimates. This fraction is removed just prior to snowmelt. Recommended default is 0.15 (Hood et al 1999). Value:  $0.0 \leq \text{decimal} \leq 1.0$ . Units: proportion. Default = 0.15.

## 9.12 WinterSTD (optional)

Standard deviation of winter temperatures. Used in conjunction with ColdTol to determine if cohorts survive the estimated extreme coldest temperature each month. Cohort is killed when:

$$[T_{ave} - 3 \times \text{WinterSTD}] < \text{ColdTol}$$

sensu Court 1951. Value:  $\text{decimal} \geq 0.0$ . Units: °C.

## 9.13 ClimateFileName

This parameter gives the name of the climate file for the ecoregion. The user may specify the same file for multiple ecoregions. Note that this parameter is required even if the climate library is used. However, the value(s) for this parameter will be ignored if the climate library is specified in a ClimateConfigFile (section 3).

## 10 Input File – DisturbanceReductions

This file contains parameters that define how different disturbances impact the wood, roots and foliage being added to or removed from the dead pools.

### 10.1 Example file:

```
LandisData  DisturbanceReductions
>>-----
DisturbanceReductions  fire  wind  harvest  bda
WoodReduction          0.33  0     0.70     0
FolReduction           1     0     0         0
RootReduction          0     0     0         0
DeadWoodReduction      0.7   0     0         0
LitterReduction        0.9   0     0.1       0
```

### 10.2 LandisData

This parameter's value must be "DisturbanceReductions".

### 10.3 DisturbanceReductions Table

This table contains the proportional reductions that should be applied to wood, root and foliage biomass as well as dead wood and litter when different disturbance types are applied. The first value in the table should be 'DisturbanceReductions'.

The columns of the table represent the different disturbance types. These can be any disturbance types that are in use in the model. Including disturbance types in this table that are not used in a given simulation scenario is acceptable and has no impact.

The rows of the table represent the different impacts on biomass or dead pools that disturbances might have. The five rows should be labeled: 'WoodReduction', 'FolReduction', 'RootReduction', 'DeadWoodReduction' and 'LitterReduction'. The values for WoodReduction, RootReduction and FolReduction represent the proportion of biomass (wood, root or foliage) that gets reduced prior to the transfer of biomass into the dead wood (wood and root) or litter (foliage) pools. These values represent how much of the cohorts are consumed by the disturbance (e.g., fire), or removed from the site as part of the disturbance (e.g., harvest).

The values for DeadWoodReduction and LitterReduction represent the proportion of the dead pool mass that is consumed or reduced due to the disturbance. The reductions of the dead pools are applied to the existing pool masses prior to additions caused by a new disturbance.

## 11 Input File - Output-PnET

This file contains parameters for the optional PnET-Succession output extension.

### 11.1 Example file:

```
LandisData "Output-PnET"
```

```
Timestep 10
```

```
Species All
```

```

Biomass                output/biomass/{species}/biomass-{timestep}.img
AbovegroundBiomass     output/AGbiomass/{species}AG/biomass-{timestep}.img
BelowgroundBiomass     output/BelowGroundBiom/BGB-{timestep}.img
WoodySenescence        output/WoodySenescence/{species}/WoodySen-{timestep}.img
FoliageSenescence      output/FolSenescence/{species}/FolSen-{timestep}.img
LeafAreaIndex          output/LeafAreaIndex/lai-{timestep}.img
Establishment          output/Establishments/{species}/Est-{species}-{timestep}.img
EstablishmentProbability output/EstabProb/{species}/EstProb-{species}-{timestep}.img
MonthlyNetPsn          output/MonthlyNetPsn/MonthlyNetPsn-{timestep}.img
MonthlyFolResp         output/MonthlyFolResp/MonthlyFolResp-{timestep}.img
MonthlyGrossPsn        output/MonthlyGrossPsn/MonthlyGrossPsn-{timestep}.img
MonthlyMaintResp       output/MonthlyMaintResp/MonthlyMaintResp-{timestep}.img
Water                  output/SoilWater/water-{timestep}.img
SubCanopyPAR           output/SubCanopyPar/SCpar-{timestep}.img
CohortsPerSpecies     output/CohortsPerspecies/{species}/cohorts-{timestep}.img
AnnualPsn              output/AnnualPsn/{species}/AnnualPsn-{timestep}.img
WoodyDebris            output/WoodyDebris/WoodyDebris-{timestep}.img
Litter                 output/NonWoodyDebris/Litter-{timestep}.img
AgeDistribution        output/AgeDist/Age-{timestep}.img
CohortBalance          output/TotalCohorts.csv
EstablishmentTable     output/EstablishmentTable.csv
MortalityTable         output/MortalityTable.csv

```

### 11.2 LandisData

This parameter's value must be "Output-PnET".

### 11.3 Timestep

This parameter is the time step of the extension. Value: integer > 0. Units: years.

## 11.4 Species

This keyword lists the species for which data are to be output. Value: space-delimited list of species names or the generic term All.

## 11.5 Output File Name Templates

Subsequent input lines list the variables to be output along with the output file naming rules. List one variable per line. Each line begins with the variable to be output, followed by a description of where the output files are placed and their naming convention. The first portion lists the directory where the files should be placed, relative the location of the scenario text file (e.g., output/agemaps/). The second portion includes one or two variables (depending on the variable) for creating file names. {species} will be replaced with the species name. {timestep} will be replaced with the output time step. Other characters can be inserted as desired. An appropriate file extension (e.g., img, csv) should also be included for the type of output. For example:

```
Water output/SoilWater/water-{timestep}.img
```

**Note: Most output maps are not compatible with the .gis map output type because the values are not integers.**

Here are the valid variable names and the required naming variables and file type of the output files: All variables are optional.

Biomass, path/{species}/filename-{timestep}.mapextension

Map. Units: g/m2. Biomass is the sum of wood and roots, but not foliage biomass.

AbovegroundBiomass, path/{species}/filename-{timestep}.mapextension

Map. Units: g/m2. AbovegroundBiomass includes aboveground wood and foliage.

BelowgroundBiomass, path/filename-{timestep}.mapextension

Map. Units: g/m2. BelowgroundBiomass includes roots only.

WoodySenescence, path/{species}/filename-{timestep}.mapextension

Map. Units: g/m2. Outputs woody biomass of the species added to woody dead pool.

FoliageSenescence, path/{species}/filename-{timestep}.mapextension

Map. Units: g/m2. Outputs foliage biomass of the species added to litter dead pool.

LeafAreaIndex, path/filename-{timestep}.mapextension

Map. Units: m2.

Establishment, path/{species}/filename-{timestep}.mapextension

Map. Units: # cohorts

EstablishmentProbability, path/{species}/filename-{timestep}.mapextension

Map. Units: probability.

MonthlyNetPsn, path/filename- {timestep}.mapextension

Map. Units: g/m2. \*

MonthlyFolResp, path/filename- {timestep}.mapextension

Map. Units: g/m2. \*

MonthlyGrossPsn, path/filename- {timestep}.mapextension

Map. Units: g/m2. \*

MonthlyMaintResp, path/filename- {timestep}.mapextension

Map. Units: g/m2. \*

\* NOTE: for variables above denoted MonthlyX, all outputs represent the monthly values for the last year of the timestep, including spinup.

Water, path/filename- {timestep}.mapextension

Map. Units: mm.

SubCanopyPAR, path/filename- {timestep}.mapextension

Map. Units: W/m2 or mmol/m2, depending on input units.

CohortsPerSpecies, path/ {species}/filename- {timestep}.mapextension

Map. Units: count.

AnnualPsn, path/ {species}/filename- {timestep}.mapextension

Map. Units: g/m2.

WoodyDebris, path/filename- {timestep}.mapextension

Map. Units: g/m2.

Litter, path/filename- {timestep}.mapextension

Map. Units: g/m2.

AgeDistribution, path/filename- {timestep}.mapextension

Map. Units: # cohorts

CohortBalance, path/filename.csv

Comma-delimited text file containing landscape total or average values of the following variables for each time step.

# of cohorts (landscape total)

Average Age of all cohorts on the landscape (years)

Average Biomass / site (g/m2)

Average LAI / site (m2)

Average Water / site (mm)

SubCanopy PAR / site (W/m2 or  $\mu\text{mol}/\text{m}^2$ , depending on input units)

Litter / site (gDW/m2)

WoodyDebris / site (gDW/m2)



AverageBelowGround / site (g/m<sup>2</sup>)

AverageFoliage / site (g/m<sup>2</sup>)

AverageNSC / site (gC/m<sup>2</sup>)

AverageAET / site (mm/yr)

EstablishmentTable, path/filename.csv

Comma-delimited text file containing landscape total number of cohorts established (by species and cause) for each time step.

MortalityTable, path/filename.csv

Comma-delimited text file containing landscape total number of cohorts killed (by species and cause) for each time step.

## 12 Input File – PNEToutputsites

This file contains parameters for the site data output extension. This extension outputs state variables each month for individual sites. This extension is used primarily for calibrating input parameters because it slows model execution.

### 12.1 Example file:

```
LandisData      PNEToutputsites

>>PNEToutputsites      MapCoordinatesX      MapCoordinatesY      MapCoordinatesMaxX      MapCoordinatesMaxY
>>-----
>>Site1 715,187.037 4,413,258.694      734284.375 4413934

PNEToutputsites      Row      Column
>>-----
Site1 1 1
Site2 1 2
```

### 12.2 LandisData

This parameter's value must be "PNEToutputsites".

### 12.3 PNEToutputsites

This keyword is followed by either map coordinate keywords or grid coordinate keywords (row/column). Each site (cell) to be output is listed on subsequent lines with a Site number and the appropriate coordinates (map coordinates must be compatible with the input maps). See example above for syntax.

## 13 Output file - SiteData Table (Optional PNEToutputsites output)

This comma-delimited table contains site-level PnET state variable values at the end of each month from the start of the spin-up period to the end of the simulation. The sites reported are specified in the input file. Values are for the entire cell and include the presence of all species and cohorts on the cell. Units for each variable are given in the header. This output is turned on in the PnET-Succession Input File by specifying the cell(s) to be output.

### 13.1 Time

Simulation year.

### 13.2 Ecoregion

Ecoregion for the cell.

### 13.3 SoilType

Soil type assigned to the cell's ecoregion.

### 13.4NrOfCohorts

Number of cohorts (all species) occurring on the cell.

### 13.5MaxLayerStdev

Maximum standard deviation of biomass of all cohorts present on the cell. Used to calculate the number of canopy layers (section 2.4).

### 13.6Layers

Number of canopy layers on the cell.

### 13.7PAR0

Photosynthetically Active Radiation (light) above the upper canopy layer. Same units as PAR in the input climate file.

### 13.8Tmin(C)

Tmin from the climate file.

### 13.9Tave(C)

Mean air temperature (°C), computed as the average of TMin and TMax from the climate file.

### 13.10 Tday(C)

Mean air temperature (°C) in the daytime, derived as the average of Tave and TMax.

### 13.11 Tmax(C)

Tmax from the climate file.

### 13.12 Precip(mm\_mo)

The monthly precipitation (as read from the climate file, mm/mo).

### 13.13 CO2(ppm)

Monthly CO<sub>2</sub> concentration (as read from the climate file, parts per million)

### 13.14 O3(cum\_ppb\_h)

Monthly ozone dosage (as read from the climate file). Units should be ppb-h (see 6.3.8).

### 13.15 RunOff(mm\_mo)

Monthly runoff that occurs from precipitation when the soil is saturated (mm/mo).

### 13.16 Leakage(mm)

Water lost out of the bottom of the rooting zone.

### 13.17 PET(mm)

Potential EvapoTranspiration. Potential evaporation is calculated as a simplified Penmann-Monteith calculation according to Stewart and Rouse (1976) as presented in Cabrera et al. (2016).

### 13.18 Evaporation(mm)

Precipitation lost to evaporation from the soil surface as a function of the LAI on the site.

### 13.19 Transpiration(mm)

Transpiration of all cohorts.

### 13.20 Interception(mm)

Precipitation intercepted by foliage and stems and not entering the soil.

### 13.21 PrecLoss(mm/mo)

Monthly precipitation runoff that occurs due to surface conditions (e.g., slope, impervious surface) when the soil is not fully saturated (mm/mo).

### 13.22 Water(mm/m)

Proportional amount of soil water as calculated by the bulk hydrology model (mm/m). Note that this gives the amount of water at the end of the month. To compute the amount of water at the beginning of the month, sum water, evaporation and transpiration.

### 13.23 PressureHead(m)

Pressure head (at the end of the month) as calculated by the bulk hydrology model (m).

### 13.24 Available water (mm)

Amount of available soil water, above the soil frost line (if any). Available water is  $\text{Water} * \text{RootingDepth}$  multiplied by the proportion of the soil that is above the frost line.

### 13.25 SnowPack(mm)

Water equivalent contained in the snowpack (mm).

### 13.26 LAI(m2)

Leaf Area Index (all species combined)

**13.27 VPD(kPa)**

Mean vapor pressure deficit for the month (kPa).

**13.28 GrossPsn(gC/mo)**

Gross photosynthesis of all species combined (gC/ m<sup>2</sup>/mo).

**13.29 NetPsn(gC/mo)**

Net photosynthesis of all species combined (gC/m<sup>2</sup>/mo).

**13.30 MaintenanceRespiration(gC/mo)**

Maintenance respiration of all species combined (gC/mo).

**13.31 Wood(gDW)**

Sum of aboveground woody biomass of all species (gDW).

**13.32 Root(gDW)**

Sum of root biomass of all species (gDW)

**13.33 Fol(gDW)**

Sum of foliage biomass of all species (gDW).

**13.34 NSC(gC)**

Sum of NSC (Non-structural carbon) of all species (gC).

**13.35 HeteroResp(gC\_mo)**

Heterotrophic respiration (decay of dead pools).

**13.36 Litter(gDW/m<sup>2</sup>)**

Biomass (all species) in the litter dead biomass pool (gDW/m<sup>2</sup>).

**13.37 CWD(gDW/m<sup>2</sup>)**

Biomass (all species) in the coarse woody debris dead biomass pool (gDW/m<sup>2</sup>).

**13.38 WoodySenescence (gDW/m<sup>2</sup>)**

Total woody biomass added to the dead pool each month.

**13.39 FoliageSenescence (gDW/m<sup>2</sup>)**

Total litter biomass added to the dead pool each month.

### 13.40 SubcanopyPAR

Photosynthetically Active Radiation (light) below all canopy layers (i.e., at ground level). Same units as PAR in the input climate file.

### 13.41 SoilDiffusivity(mm2\_s)

Thermal diffusivity of the soil given the texture and water content.

### 13.42 FrostDepth(mm)

Depth to the soil ice layer.

### 13.43 LeakageFrac(-)

Computed LeakageFrac (dependent on frost depth).

## 14 Output file - CohortData Table (Optional PNEToutputsites output)

This table contains monthly PnET cohort-level state variable values for the sites specified in the input file. A file is created when a cohort is established, and the records give month-end state variable values for the cohort from establishment to death (or the end of the simulation). Files are also produced for cohorts established during the spin-up period. Units for each variable are given in the header. This output is turned on in the PNEToutputsites input file by specifying the cell(s) to be output.

### 14.1 Time(yr)

Simulation year.

### 14.2 Age(yr)

Current age of the cohort (calendar years).

### 14.3 TopLayer(-)

The highest layer number to which the cohort is assigned, with 0 being the lowest layer.

### 14.4 LAI(m2)

Leaf area index of the cohort.

### 14.5 GrossPsn(gC/m2/mo)

Cohort gross photosynthesis (gC/m2/mo).

#### 14.6FolResp(gC/m<sup>2</sup>/mo)

Cohort foliar respiration (gC/m<sup>2</sup>/mo).

#### 14.7MaintResp(gC/m<sup>2</sup>/mo)

Cohort maintenance respiration, including tissue repair and nutrient transport (gC/m<sup>2</sup>/mo). This amount comes out of the NSC pool.

#### 14.8NetPsn(gC/m<sup>2</sup>/mo)

Cohort net photosynthesis (gC/m<sup>2</sup>/mo).

#### 14.9Transpiration(mm/mo)

Cohort water actually lost to transpiration (mm/mo).

#### 14.10 WUE(g/mm)

Cohort mean water use efficiency (gC/mm H<sub>2</sub>O).

#### 14.11 Fol(gDW/m<sup>2</sup>)

Biomass of the cohort foliage pool (gDW/m<sup>2</sup>).

#### 14.12 Root(gDW/m<sup>2</sup>)

Biomass of the cohort root pool (gDW/m<sup>2</sup>).

#### 14.13 Wood(gDW/m<sup>2</sup>)

Biomass of the cohort wood pool (gDW/m<sup>2</sup>).

#### 14.14 NSC(gC/m<sup>2</sup>)

Amount of carbon in the cohort non-structural carbon pool (gC/m<sup>2</sup>).

#### 14.15 NSCfrac(-)

Fraction of carbon in the cohort non-structural carbon pool relative to active biomass (NSC / (FActiveBiom \* (wood + root+ foliage). Cohorts die when NSCfrac is <0.01 at the end of a calendar year.

#### 14.16 fWater(-)

Reduction factor related to water availability.

#### 14.17 water(mm/m)

Average volumetric soil water value used to compute fWater across all cohort sublayers.

### 14.18 PressureHead(mm)

Pressure head value used to compute fWater.

### 14.19 fRad(-)

Reduction factor related to light availability at the top of the canopy layer occupied by a cohort.

### 14.20 fOzone(-)

Reduction factor related to ozone effects.

### 14.21 DelAmax(-)

Enhancement factor related to CO<sub>2</sub> effects on photosynthesis.

### 14.22 fTemp\_psn(-)

Reduction factor related to sub-optimal temperature for photosynthesis.

### 14.23 fTemp\_resp(-)

Reduction factor related to temperature effects on maintenance respiration.

### 14.24 fAge(-)

Reduction factor for age-related declines in photosynthesis efficiency.

### 14.25 LeafOn(-)

Indicates growing season status. When TRUE, new foliage can be added and old foliage has not yet been dropped.

### 14.26 FActiveBiomass(gDW\_gDW)

Fraction of active biomass. Indicates the computed fraction of wood biomass that is considered active and able to transport water to support foliage.

### 14.27 AdjFolN(gN\_gC)

Adjusted foliar nitrogen (gN/gC). When using the optional FolNInt and FolNSlope parameters, the adjusted FolN can be different from the species FolN value, depending on canopy position and the resulting PAR. Averaged across the cohort's subcanopy layers.

### 14.28 AdjFracFol(-)

Adjusted FracFol. When using the optional MaxFracFol and FracFolShape parameters, the adjusted FracFol can be different from the species FracFol value, depending on canopy position and the resulting PAR. Averaged across the cohort's subcanopy layers.



### 14.29 CiModifier(-)

Reduction factor for gas conductance due to stomatal closure caused by water stress and modified by ozone. Value of 1.0 indicates no reduction in conductance. Averaged across the cohort's subcanopy layers.

### 14.30 AdjHalfSat(-)

Adjusted HalfSat. When using the optional CO2HalfSatEff, the adjusted HalfSat can be different from the species HalfSat value, depending on concentration of CO<sub>2</sub>. Averaged across the cohort's subcanopy layers. Units are typically  $\mu\text{mol}/\text{m}^2/\text{sec}$  or  $\text{W}/\text{m}^2$  and will be the same as the input units of PAR in the climate input file and the species HalfSat value.

### 14.31 Limiting Factor

Indicates the most limiting factor on photosynthesis, including death by cold (and the temperature that killed the cohort).

## 15 Output file – Establishment Table (Optional PNEToutputsites output)

This comma-delimited table reports site-level establishment information for each species. The reported values reflect state variables in the model at intervals of one PnET-Succession time step.

### 15.1 Year

Simulation year (timestep).

### 15.2 Species

Species.

### 15.3 Pest

The average probability of establishment for the species during the given time step as a function of the average monthly values of fWater and fRad. The average probability is:

$$P_{est} = fRad\_Avg \times fWater\_Avg \times MaxPest$$

### 15.4 FWater\_Avg

The average water availability reduction factor (fWater) value across all growing season months in the time step.

### 15.5FRad\_Avg

The average light availability reduction factor (fRad) value across all growing season months in the time step.

### 15.6ActiveMonths

The number of growing season months within the timestep.

### 15.7Est

Indicates if an establishment of the species by seeding would occur on the site in the time step, if seeds are able to reach the site and the site did not have planting, serotiny or resprouting occur. Actual establishment additionally requires a source tree within seeding distance.

## 16Output file – Mortality Table (Optional PNEToutputsites output)

For each time step the model outputs the number of cohorts of each species killed by the major sources of mortality (Succession (competitive exclusion), Harvest, Fire, Wind and Other).

## 17Output file – Establish Table (Optional PNEToutputsites output)

For each time step the model outputs the number of cohorts of each species established by the four sources of cohort establishment (Planting, Serotiny, Resprouting and Seeding).

## 18Appendix. Calibration tips.

### Calibration tips

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#### General

One of the compelling features of PnET-Succession is that its parameters are mostly empirically estimable values, and it autonomously produces very realistic growth responses under a wide variety of abiotic conditions when its parameters are set correctly. The developers of the PnET model claim (perhaps too optimistically) that PnET does not need calibration, but there are some changes that have been made to PnET-Succession to make it tractable at landscape scales that cause the model to require calibration. However, of the dozens of parameters used by PnET-Succession, only a few typically need calibration, and most applications can use default values for most parameters.

I calibrate each species separately, using a single cohort initialized on a single cell and grown for about 150 years. I compare simulated biomass (wood) growth through time with empirical biomass growth curves, or those generated by the forest stand model FVS. Use the PNEToutputsites option to produce the cohort growth output files needed for calibration.

To produce predictable competitive interactions that can be controlled by intuitive modification of a small number of parameters for individual species, it is best to use common parameter values across all species as much as possible. These can be conveniently maintained (and modified as necessary) in the GenericPnETSpeciesParameters file. Other parameters vary by life history trait or growth form, and each species having a particular life history trait should have the same (or very similar) parameter value to represent that trait. Examples of such traits include shade tolerance (HalfSat), drought tolerance (H3 & H4), extinction coefficient (k), relationship between foliar N and photosynthetic capacity (AmaxA and AmaxB), leaf weight change by canopy position (SLWdel) and fraction of foliar biomass relative to active wood biomass (FracFol). Parameters that the model is highly sensitive to (for which common values should be used as much as possible) include: MaintResp, TOroot, TOwood, BFolResp, FracBelowG, FracFol and FrActWd.

Calibration is ideally done by modifying only FolN (to control relative productivity), FracFol (to control gross LAI and the initial growth part of the growth curve), SLWmax (to fine-tune mature LAI and the ability to compete for light). FracFol and FrActWd can have a profound effect on the competitive ability of cohorts, so to achieve reasonable competitive interactions it is best to find a common value to use for both of these parameters for all species having a common growth form (deciduous v. conifer), modifying them only for species that have exceptionally unusual growth curves. FracFol must generally be much higher for conifers because their AmaxA and AmaxB terms produce much less growth per unit of foliage. Find a value that generally works for all species of a growth form and be hesitant to modify it for individual species unless there is a compelling biological or empirical reason. FrActWd controls the amount of foliage in mature cohorts and the maximum height of the growth

curve, and should vary from the common value mostly when the height of the growth curve needs to be suppressed (e.g., shrub species).

I set up a landscape with a single active cell, and initialize it with a single species at a time. I calibrate species under good to ideal growing conditions, recognizing that their growth will be less under poorer conditions and when competing because stress factors will be greater. However, if you know the conditions under which your growth curves were generated, use those conditions. I calibrate on a soil with relatively high water holding capacity (e.g., SALO) and a weather stream with average or better precipitation for the regions where the species are typically found, and calibrate to the higher range of the growth curves available. The key is to calibrate under nearly ideal conditions, or conditions that reflect those that produced the growth curves to which you are calibrating. I use a constant climate for calibrating, to avoid confounding the calibration by extreme events. You should use growing season temperatures approximating  $PsnTOpt$ . This is most easily done by temporarily setting  $PsnTxxx$  values to match those of the climate inputs. Note that the model compares  $PsnTxxx$  values to  $Tday$  (output in the Site.csv file), not  $Tmax$ . Don't forget to set  $PsnTxxx$  values back to actual values when finished calibrating. I am content if my calibrated curve produces growth somewhat better than the average of my empirical growth data or near the highest growth curve, because calibration represents near ideal conditions. Use  $CO_2$  (and  $O_3$  if applicable) values typical of those experienced by the trees measured for your empirical biomass growth curves.

The calibration process is expedited by plotting simulated wood biomass against empirical growth data in a spreadsheet. I create a separate spreadsheet tab for each species and also plot foliage biomass and LAI. Other variables for which empirical data are available can also be plotted against each other. Using a macro, I can quickly copy and paste data from the cohort output file of PnET-Succession to iteratively achieve the desired behavior through multiple model runs.

MaintResp is a critical parameter for PnET-Succession because it is the primary determinant of cohort growth limitations and death, and growth is highly sensitive to variation in this parameter. Unfortunately, this parameter is rarely known empirically in the units used by the model, so it must be calibrated. If MaintResp is too low, cohorts will never die from stress and growth will be virtually unconstrained. If MaintResp is too high, growth will quickly level off because of high maintenance costs and cohorts will be highly susceptible to stress. For most studies, competitive outcomes should be a function of competition for light and water, not from a difference in maintenance costs. Therefore, it is recommended to find a value for MaintResp that works across life history traits and growth forms. This is fairly easily done. Choose some representative species having generic growth parameters (including great longevity) and simulate their growth curves, searching for a value of MaintResp that flattens (levels off) the growth curve after a few decades. Then gradually reduce MaintResp until the growth curve takes the shape of that typically seen in the growth curves of your long-lived species. Select a value of MaintResp that produces acceptable behavior for all life forms, etc., and use it for all species. In summary, you are searching for a value that is as high as it can be without creating a flattened growth curve.

Useful empirical data needed for calibration.

1. The range of FolN for each species across sites, age, canopy position. This is a great starting point and helps put bounds on acceptable values, but it is important to recognize that productivity is closely related to FolN in PnET, so it is more important that FolN reflect relative productivity among species rather than matching empirical values. That said, it is usually possible to select a value of FolN that is within the range of empirically measured values. If you have empirical values of Amax, use your values of AmaxA and AmaxB (or see Fig. 2 in Aber et al 1996) to estimate the FolN that produces that Amax value. NOTE: if you plan to implement dynamic FolN (FolNInt and FolNSlope set to values other than default), then be sure to set FolNInt and FolNSlope to the values you will use in your simulations.
2. Species SLWmax helps determine LAI. Thicker leaves (higher SLW) reduce LAI. LAI determines the ability of a cohort to compete for light, so it is important to calibrate so that LAI approximates empirical values. Again, it is more important to get LAI approximately right than to keep SLWmax within the empirical range. Note that higher LAI increases within canopy competition for light and will modestly depress growth even of a single cohort.
3. Growth curves by species. Growth and yield tables are useful for this purpose and volume measures can be converted to biomass using specific gravity values for the species (Miles and Smith 2009). Ensure that units are the same as output by PnET-Succession. Most growth and yield tables account for only the merchantable part of trees, so consider them to be near the lower boundary of the woody biomass output of PnET-Succession. I have recently begun calibrating to growth curves generated by the FVS model for various soil types. One advantage is that the curves for all species are directly comparable, and your parameters will produce realistic competition among species. The primary feature of the growth curve that you should calibrate to is the maximum height of the curve. The rate of initial increase can be matched quite readily using FracFol, but if you have different values of FracFol across species, the lower valued species will compete very poorly. So, avoid the temptation to match the initial rate of increase and focus instead on the height of the curve. Make modest exceptions for species that tend to outgrow their competitors when establishing on open sites (e.g. pioneer species) and for species that have unusually slow growth compared to competitors with similar life history traits (e.g., white cedar, hemlock). I generally use slightly higher FracFol values for pioneer species and slightly lower values for shade tolerant species. The shape of the senescence decline can be controlled by PsnAgeRed, but again resist the temptation to calibrate this individually for each species unless the species has well-known anomalous senescence characteristics.

### Calibration procedure

1. First, set parameter values that are known from the literature (e.g., TOFol, FolN, SLW, HalfSat, PsnTMin, PsnTOpt, k, etc.). When there is considerable range in empirical values, begin with an intermediate value. HalfSat should be set to reflect shade tolerance, because the LANDIS-II shade tolerance parameter has no effect in PnET-Succession. Some PnET parameters (e.g., SLWDel, AmaxA, AmaxB, Q10, DVPD1, DVPD2) are hard to estimate and most studies use generic values (e.g., Aber et al 1995,

- Ollinger and Smith 2005). Decide whether you will use the Wythers=TRUE option; I recommend using it. NOTE: if you use dynamic FolN, you should start with FolN values near the bottom of their empirical range.
2. Use the GenericPnETSpeciesParameters file for all parameters that are not varied among species. Set parameters that will be held constant for your particular experiment or study (e.g., TOrroot/wood, BaseFolResp, InitialNSC, etc.) Again, it is advisable to hold as many parameters constant across species as possible to allow you less confounded control of competitive interactions with the remaining parameters.
  3. Temperature parameters provide an important link between climate and species growth, and they should be calibrated carefully when conducting climate-related studies. LeafOnMinT controls the length of the growing season by comparing LeafOnMinT to monthly Tmin. You should verify that the appropriate months are active (LeafOn=TRUE) given the climate inputs. LeafOnMinT should vary coarsely according to leaf-on and leaf senescence dates, and typically ranges between 2-4 °C for temperate species.
  4. PsnTMin controls how productive the species is at the beginning and end of the growing season, and will produce a growth response to shifts in temperature even when the number of growing season months is unchanged. Similarly, in some years you may get some modeled photosynthesis activity in a month when the species is not typically active for the entire month, but this is not a problem as long as NetPsn is quite low in those months. Note that PsnTMin, PsnTOpt and PsnTMax values are compared to Tday (average daytime temperature), not Tmax (daytime high temperature), and should be set accordingly. I typically vary PsnTMin among species relative to the isotherms at their northern range boundaries.
  5. In lieu of empirical values, PsnTOpt can be estimated using the average mid-summer temperature at the center of the species' range. .
  6. For the most realistic high temperature response, use the DTEMP=true option. Empirical values for maximum temperature limits to photosynthesis are typically instantaneous photosynthesis rates, while PnET-Succession interprets PsnTMax as the average daily temperature above which there is no net photosynthesis. Thus, PsnTMax should be lower than the typical bio-chemical limit of 38-39 °C. I use PsnTMax to make more northerly species less competitive than southerly species as climate warms. I typically vary PsnTMax among species relative to the isotherms at their southern range boundaries unless those boundaries are clearly related to a factor other than temperature.
  7. Calibration tuning is best done by matching simulated wood biomass increase to empirical biomass values for a species through time. If your empirical values represent something less than whole tree aboveground biomass, your tuned values should be higher than the empirical values. Your goal is not to match the empirical curve exactly, but to approximate it. Simulate a monoculture and plot Wood and Root+Wood (whole tree) biomass through time. There are some published estimates of whole tree biomass through time by species groups to provide some indication of belowground biomass (e.g., Smith et al 2006). Comparison of empirical aboveground and whole tree biomass can help you estimate the FracBelowG parameter. The calibration simulations should run for at least as many years as your empirical data, and it may be informative to see how the model extrapolates growth beyond your empirical data. When you find it difficult to match an empirical growth curve without setting a parameter outside its empirical range

or to a very different value than similar species, modifying FracBelowG (or TOWood or TORoot) can produce a marked effect on the growth curve with a small change in value. An increase in one pool will reduce the other pool. It would be beneficial to have some justification for your selected value from empirical evidence.

8. Choose a single soil type for calibration of all species, preferably a soil type that is both common and productive. This enables you to estimate comparable life history trait parameters, ensuring that competition is realistic. Ultimately, growth is limited by water, so you want to calibrate under somewhat ideal conditions to reflect realistic competition for water. I typically do not choose the soil type with the greatest water holding capacity, but one that is above average. Soil water is determined by inputs (precipitation) and outputs (PrecLossFrac, percolation out of the rooting zone (controlled by SoilType) and transpiration). Tuning of soil water is done primarily with PrecLossFrac, LeakageFrac and SoilType, which assumes that transpiration is correct if the photosynthesis and growth behavior is correct. PrecLossFrac typically represents water lost where slope is sufficient for water to run off before it can enter the soil. You should calibrate for a slope condition that is flatter than average in your study area. Wetland ecoregions should have ecoregion parameters (RunoffCapture and LeakageFrac) that produce appropriate soil saturation. RunoffCapture represents the maximum depth of standing water across the ecoregion, and LeakageFrac can be used to reflect any impermeable soil layer that reduces drainage. In lieu of empirical data to help set these parameters, tune them such that the species typically found there can survive and the hydrology mimics reality.
9. Calibrate all species under optimal temperature and precipitation conditions, which will result in growth reductions under other conditions. If you don't uniformly calibrate species under optimal conditions, then their competition will be unrealistic and unpredictable during simulations. This is most easily done by using a fixed annual weather stream (long-term monthly averages), using typical monthly precipitation values.
10. You will need starting parameter values. Here are some generic ones for temperate forest species that are widely used in PnET-Succession simulations:
  - a. SLWDel: 0.0 for evergreen, 0.2 for deciduous
  - b. AmaxA: 5.3 for evergreen, -46 for deciduous
  - c. AmaxB: 21.5 for evergreen, 71.9 for deciduous
  - d. BFolResp: 0.1
  - e. k: 0.5 for evergreen, 0.58 for deciduous
  - f. DNSC: 0.05
  - g. Q10: 2
  - h. DVPD1: 0.05
  - i. DVPD2: 2
  - j. IMAX: 5-10 (there is a significant performance cost with values higher than these)
  - k. TOWood: 0.01
  - l. TORoot: 0.02

Here are some other values I have used that you may need to modify for your species:

- a. HalfSat: 300 ( $\mu\text{mol/s}$ ) for shade-intolerant, 100 for shade-tolerant; I use a gradient across 5 classes.
- b. H1.H2: 0/4 for most species. Use -3.3/2 for species that are very waterlogging tolerant. This allows some photosynthesis even on saturated soils. Use the PnET-

- Succession function worksheet to find values that produce the desired reduction in photosynthesis on saturated soil.
- c. H3/H4: 100/140 for drought intolerant, 118/153 for drought tolerant; I use a gradient across 4 classes.
  - d. MaintResp: 0.002
  - e. PsnAgeRed: 5
  - f. FracBelowG: 0.33. You may wish to systematically vary this fraction to reflect life history strategy: slightly lower (0.31) for shade-intolerant species and higher (0.35) for shade-tolerants
  - g. FracFol: 0.10 for evergreen, 0.03 for deciduous. You may wish to systematically vary this fraction to reflect life history strategy: slightly upward for pioneer species and downward for slow-growing species.
  - h. FracActWd: 0.00004 (shrubs may require values of 0.0001-0.0002).
  - i. PrecipEvents: 9-11
11. Relative growth rate among species should be controlled with FolN. Conifers and deciduous species must have FolN scaled separately because they use different values of AmaxA and AmaxB. Unless specific, high quality estimates of AmaxA and AmaxB are known, it is recommended to use the values commonly used in PnET publications (see above). Published values of FolN are commonly available, but use these only as a starting point. **In PnET-Succession, FolN is the primary parameter that controls relative growth rate (Amax) among species.** Set FolN of your suite of species based on what is known about relative growth rates among your species. Be aware that differences in shade tolerance and drought tolerance will prevent FolN from scaling perfectly with growth rate among all species. I have found that FolN should be increased ~0.1 for each decrease in shade tolerance class (of 5 classes, binned according to HalfSat) to produce approximately the same growth rate. However, you should be able to use a FolN value for each species that is within the range of empirical values. If you find you must use a FolN value that is out of range, modify other parameters to bring it within range, starting with FracFol, FracActWd or FracBelowG. Don't forget that you will have a different range of FolN values for species with different AmaxA and AmaxB values. If you are using dynamic FolN, check the AdjFolN values to see if they vary within empirical limits.
12. Foliage biomass is controlled by FracFol and FrActWd, and you will find that FracFol also tightly controls the lag time of significant biomass increase (early years). Because this lag time has a strong effect on competitive ability in PnET-Succession, it is recommended that you attempt to use common (or very similar) values of FracFol and FrActWd. It may be preferable to have similar lag times than to match empirical lag times closely; this will produce better competitive interactions in the model. Super slow-growing species (e.g., white cedar, black spruce) may be an exception. Often empirical growth curves are measured in mixed forests, and shade-tolerant species (e.g., eastern hemlock) will tend to have longer lag times than when grown in the open as in the calibration simulations. You should monitor the shape of the foliage curve along with the biomass curve. If FrActWd is too high, foliage biomass will decline markedly with age, which is not likely realistic. If it is too low, foliage biomass will increase indefinitely, which is also not realistic. Experience shows that a value of 0.00004 will produce a level or slightly declining foliar biomass after about 50 years.



13. Before adjusting the height of the biomass curve, verify that LAI is not way out of bounds for the species. LAI is controlled by FracFol, FrActWd, SLWmax (and SLWdel). FracFol directly controls the amount of foliar biomass, and LAI is calculated by dividing foliage biomass by SLWmax. Note that FolN and SLWMax are generally inversely correlated in the literature. SLWdel controls the reduction of SLW from top to bottom in the canopy, and can have a large effect on LAI when there is a lot of foliage biomass. To estimate whether SLW at the bottom of the canopy may be reasonable, use  $SLWmin = SLWmax - SLWdel * \text{foliage biomass}$ . FrActWd controls the amount of foliage biomass later in life, and has an important effect on maximum LAI achieved; use this to control LAI if SLWmax values are well established empirically. In lieu of empirical values, LAI should generally range between 2-4 for shade intolerant species and 4-6 for shade tolerant species, with values over 7 not unreasonable for highly shade tolerant species. LAI is fairly insensitive to changes in FolN, so fine tuning FolN will not spoil your tuning of LAI.
14. The height of the biomass growth curve is primarily controlled by FolN, FracFol, FrActWd, but is ultimately determined by water limitation. Remember that FracBelowG, TORoot, TOWood also have an effect. H3/H4 has a slight effect by modifying water stress and HalfSat can modify light stress. Also remember that MaintResp can cause the curve to plateau regardless of other parameter settings. It is recommended to make major adjustments to the height of the biomass curve with FracFol, FrActWd and then fine-tune it using FolN and SLWmax (within empirical limits). Note that raising LAI using SLWmax generally reduces the height of the growth curve slightly, because less light penetrates deeper into the canopy. When calibrating conifers, note that higher values of TOfol depress the growth curve because more of the NetPsn is needed to build leaves. Thus, higher FolN may be needed for species with shorter leaf longevity compared to similar species with longer leaf longevity.
15. The timing of the peak of the biomass growth curve is primarily determined by species longevity. To modify the timing of the onset of senescence, adjust PsnAgeRed. A value of 5 seems to work well for most species.
16. You should be able to control most aspects of the shape of the biomass growth curve. The initial increase is controlled with FracFol, the height by FolN and FrActWd, and the decline by longevity with PsnAgeRed. Fine-tuning is done with FolN and SLW. FolN has a predictable effect on the growth curve, but you may find SLW a bit more erratic because it interacts with foliage biomass and leaf area to affect light competition within the canopy. FolN tends to shift all parts of the growth curve up or down without changing its shape. For species that accumulate markedly less biomass than most species, reduce FracFol modestly and/or increase FrActWd.
17. NetPsn is primarily controlled by FracF and FracActWd, given FolN, SLW and BaseFolResp. Transpiration is highly correlated with GrossPsn, scaled by WUE (which is modified by CO<sub>2</sub> and O<sub>3</sub> through their effect on Ci). Recall that GrossPsn is calculated by PnET-Succession using NetPsn and BaseFolResp, which is not intuitive.
18. DNSC is the target proportion of carbon reserves that is maintained, and it has little effect on cohort competition unless it is set so low that the cohort has minimal reserves to survive stress, or so high that the species can rarely be stressed enough to die. Set this in the middle of the range of empirical measures of total % sugars and starch in active tissues. In lieu of empirical values, use 0.05.

19. InitialNSC similarly has little effect on cohort competition unless values among species vary by more than an order of magnitude.
20. Establishment probability (Pest) is a function of the light and water photosynthesis reduction factors for the species at the time of establishment. The relationship with the light reduction factor is shown in Section 1.3.1 of the PnET-Succession v4.0 user guide. The two reduction factors are rarely scaled the same in that fWater commonly takes values of 1.0, while fRad may never exceed values considerably lower than that. To give both reduction factors the same weight, set each establishment modifier equal to the highest expected reduction factor value. This can be estimated by using the maximum value for each factor seen in the calibration files for each species. Typically, EstMoist will equal 1.0, and EstRad may range from 0.8-0.94. Furthermore, establishment rates will tend to be extremely high using Pest as calculated directly from the two main reduction factors. Consider the parameter MaxPest to be a tuning knob to dial all establishment rates up or down. This cannot be done on a single cell, and must be done on a landscape before you conduct landscape-scale experiments. I don't have enough experience to provide concrete guidance. I have used a management and disturbance scenario that reflects recent history and look to see that the total number of cohorts reaches an equilibrium that is not wildly different (e.g. 2x increase) than the starting number of cohorts. Because initial number of cohorts is highly uncertain depending on how the initial conditions were estimated, it may also be useful to use PnetOutputSites for a number of typical sites and see if you get a lot of cohorts established that are unable to thrive. I suggest starting with a MaxPest value of 0.4 (when time step = 10) and tune up or down from there. MaxPest is dependent on the PnET-Succession time step, meaning that you must reset its value when you change the time step. We think it may be as simple as reducing the value in proportion to the change in time step length. For example, if the length of the time step is reduced by half, also reduce the value of MaxPest by half. We are working on a way to modify the code to make MaxPest time-step invariant, but have not yet found a robust method.
21. WinterSTD. See map in Court (1951) for values in the US.
22. After several years of using PnET-Succession, I am still learning things about model behavior, and revising my calibration protocol. If you should learn something that would improve this calibration guide, please share it with me at [eric.gustafson@usda.gov](mailto:eric.gustafson@usda.gov).

**General notes.** 1) Again, to ensure realistic competition, it is advisable to use common parameter values across species, unless you have empirical data that are comparable and reliable across species. If you are planning to experimentally vary some parameters, holding the others constant will improve the signal from your experiment. Minimizing species differences in parameters such as SLWDel, PsnAgeRed, k, MaintResp, DNSC, FracBelowG, FracFol and FracActWd will make competitive interactions more predictable. Hold these as close to each other as possible, varying other parameters to calibrate as much as possible within empirical limits. However, when you cannot calibrate adequately using the common parameter values, do not hesitate to vary the one or two other parameters that will produce good performance. It is very likely that such modifications reflect biological reality. 2) Use the PnET-Succession function worksheet to help you understand how the parameters determine the behavior of cohort state variables as a function of the abiotic inputs, both intermediate variables and the ones that ultimately reflect competition and growth (e.g.,

NetPsn, foliar and wood biomass, NSCfrac). It is available from the PnET-Succession page of the LANDIS-II website ([www.landis-ii.org](http://www.landis-ii.org)). 3) It is highly recommended that you verify your calibrations by simulating several similar species together to ensure that they in fact compete as expected. If not, often a slight tweak of one parameter can bring them in line with expectation. The likely parameters to tweak are FolN (general growth rate), H3/H4 (competition for water) and HalfSat (competition for light), or FracFol (amount of foliage).

Here are some commonly used generic values, for your reference.

LandisData PnETGenericParameters

PnETGenericParameters	Value	
BFolResp	0.1	
TOroot	0.02	
TOwood	0.01	
MaxDevLyrAv	10000	
MaxCanopyLayers	3	
IMAX	5	
DNSC	0.05	<<target NSCfraction
MaintResp	0.002	
FracBelowG	0.33	
KWLit	0.1	<<0.075 for hardwoods and 0.125 for softwoods (Mattson 1987)
FolLignin	0.2	<<Range in Melillo et al (1982) = 0-0.25
Wythers	true	
PrecipEvents	11	
DTEMP	true	
RunoffCapture	0	<< upland ecoregions; may be 75 (mm) for lowlands
LeakageFrac	1.0	<<upland ecoregions; may be 0.01 for lowlands
LeakageFrostDepth	3000	
PrecIntConst	0.1	<<upland ecoregions; may be 0.05 for sparsely populated lowlands
WinterSTD	6.67	<<ranges between 6.67 (52 deg. N) and 9.2 (70N) in Siberia
MaxPest	0.4	

Here are some species parameters that have been calibrated in various ecosystems in the Midwestern U.S., mostly using a SALO soil (Gustafson et al. 2020). These can be used as a starting point, but they should be calibrated for your study area. FolN was NOT dynamic. Note that AmaxA and AmaxB are not shown, and vary between deciduous and evergreen as described above.

LandisData PnETSpeciesParameters

&gt;&gt;calibrated for Wythers=true

&gt;&gt;Upper Midwest climate adaptation study

Species	Fol N	SL W max	TO fol	Hal fSat	H1	H2	H3	H4	Leaf OnM inT	Psn TM in	Psn TO pt	Psn TM ax	Fra cFo l	Fr Act Wd	Col dT ol
abiebals	1.4	165	0.2 5	150	- 0.7 5	3	100	140	1.9	5.7	20. 5	29	0.0 9	0.0 000 4	-65
acerrubr	2.3	75	1	150	- 1.9	2.7 5	111	152	2.5	6.5	26. 5	36	0.0 29	0.0 000 4	-50
acersacc	2.3	70	1	100	0	4	105	145	2.5	6.5	24. 1	31	0.0 28	0.0 000 4	-55
betualle	2.2	70	1	150	- 1.5	2.8	105	145	2.5	6.5	21. 5	32	0.0 29	0.0 000 4	-55
betupapy	2.5	110	1	250	- 0.7 5	3	111	152	1.8	5.8	19. 5	28	0.0 31	0.0 000 4	-65
carycord	2.4	110	1	250	- 0.7 5	3	118	160	3	7	26. 3	34. 3	0.0 31	0.0 000 4	-51
fagugran	2.3	65	1	97	0	4	100	140	2.7	6.7	26. 4	34. 5	0.0 28	0.0 000 4	-50
fraxamer	2.4	80	1	200	- 0.7 5	3	105	145	2.8	6.8	26. 5	34. 5	0.0 3	0.0 000 4	-50
fraxnigr	2.4	115	1	250	- 3.3	2	105	145	2	6	23. 6	33. 3	0.0 31	0.0 000 4	-60
fraxpenn	2.4	80	1	200	- 1.5	2.8	118	160	2.5	6.5	26. 5	34. 5	0.0 3	0.0 000 4	-51
larilari	2.6	90	1	300	- 3.3	2	105	145	1.7	5.7	19. 5	30	0.0 32	0.0 000 4	-65
piceglau	1.4	210	0.2 5	200	0	4	111	152	1.6	5.7	19. 5	30	0.1	0.0 000 4	-66
picemari	1.2	210	0.2 5	200	- 3.3	2	105	145	1.6	5.6	19. 5	29	0.1	0.0 000 4	-66
pinubank	1.6	270	0.3 333	300	0	5	118	160	1.7	5.7	17. 5	28	0.1 1	0.0 000 4	-61
pinuresi	1.6	265	0.3 333	250	0	4	111	152	2.2	6.2	20. 1	30	0.1 05	0.0 000 4	-55

pinustro	1.7	215	0.5	200	-0.75	3	111	152	2.2	6.2	23	31.3	0.1	0.0004	-53
popubals	2.5	125	1	300	-0.5	4	105	145	2	5.7	20.5	29	0.032	0.0004	-65
popugran	2.5	125	1	300	-0.5	4	111	152	2.5	6.1	21	30	0.032	0.0004	-55
poputrem	2.5	125	1	300	0	4	111	152	2	6	20	29.4	0.032	0.0004	-65
prunsero	2.6	95	1	250	0	4	111	152	2.8	6.8	26.5	34.5	0.03	0.0004	-43
queralba	2.5	100	1	250	0	4	118	160	3	7	26.5	34.5	0.031	0.0004	-41
querelli	2.5	100	1	250	0	4	118	160	2.5	6.5	26	34.5	0.031	0.0004	-50
quermacr	2.5	90	1	200	0	4	118	160	2.3	6.3	26	34.6	0.03	0.0004	-53
querrubr	2.5	85	1	200	0	4	111	152	2.7	6.7	26.5	34	0.03	0.0004	-50
quervelu	2.4	85	1	200	0	4	111	152	3.1	7.1	26.5	34.3	0.03	0.0004	-41
thujocci	1.4	200	0.5	200	-3.3	2	111	152	2	6	22.8	32	0.1	0.0004	-60
tiliamer	2.3	75	1	150	0	4	105	145	2.4	6.4	24	32.5	0.029	0.0004	-51
tsugcana	1.7	110	0.333	95	0	4	100	140	2.7	6.7	26	34.2	0.08	0.0004	-65
ulmuamer	2.3	85	1	200	-1.9	2.75	105	145	2	6	26.3	34.5	0.03	0.0004	-40

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