

Equilibrium and non-equilibrium concepts in forest genetic modelling: population- and individually-based approaches

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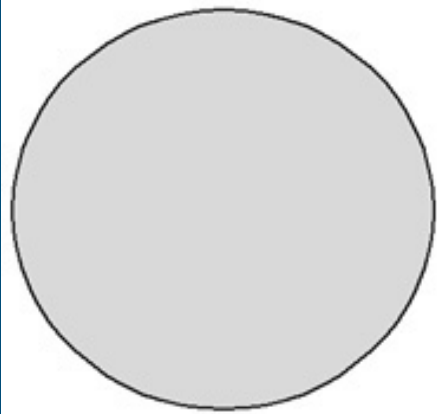
Structure of the presentation

- Equilibrium vs. non-equilibrium approaches in forest genetic modeling
 - Eq.: Population-genetic modeling
 - Non-eq.: Individually-based modeling
- Exampes non-eq modelling - traits under rapid evolution:
- Discussion
 - Pros and cons of eq. and non-eq. genetic modeling to include in regular forest models
- Conlusion
 - On which genetic model to add to existing forest model

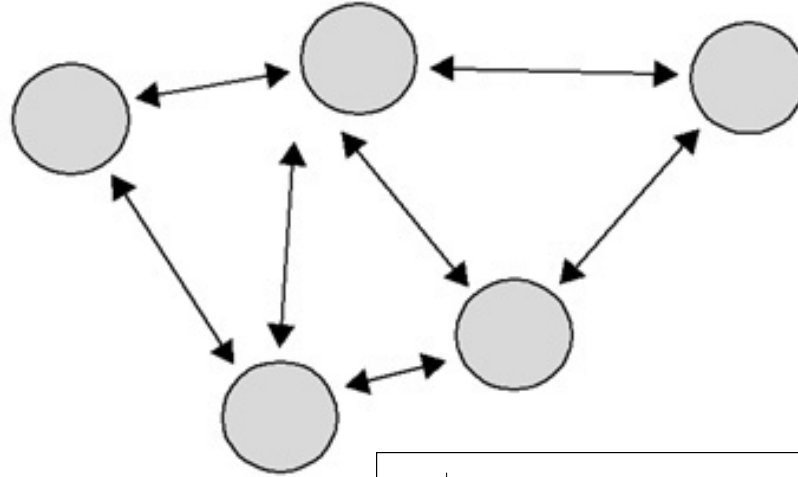
Equilibrium or demographic approach

- Assumption:
 - Environment is stationary (no trends in space nor time) relative to the rate of recovery after a perturbation
 - => following a perturbation the population returns to a previous (thus known) stable state: equilibrium
 - => we can use current knowledge on dependency of stable state to environmental factors to assess future stable states
- Traits to differentiate populations, e.g.:
 - Fecundity, survival, competition, dispersal, biomass, height, bud burst
 - i.e. usually phenotypic plastic traits (GxE interaction)
- Model parameters under study e.g.:
 - Demographic: carrying capacity (K), per capita growth rate (r)
 - Genetic: optimal phenotype (Z_{opt}), selection coefficient (ω)
- Model analyses, e.g.:
 - Recovery time (# generations) to a known (future) stable state, depending on genetic structure (dominance, epistacy) and / or spatial structure of the population
- Use:
 - Provides insight in system dynamics
 - Understanding of current patterns based on historic processes

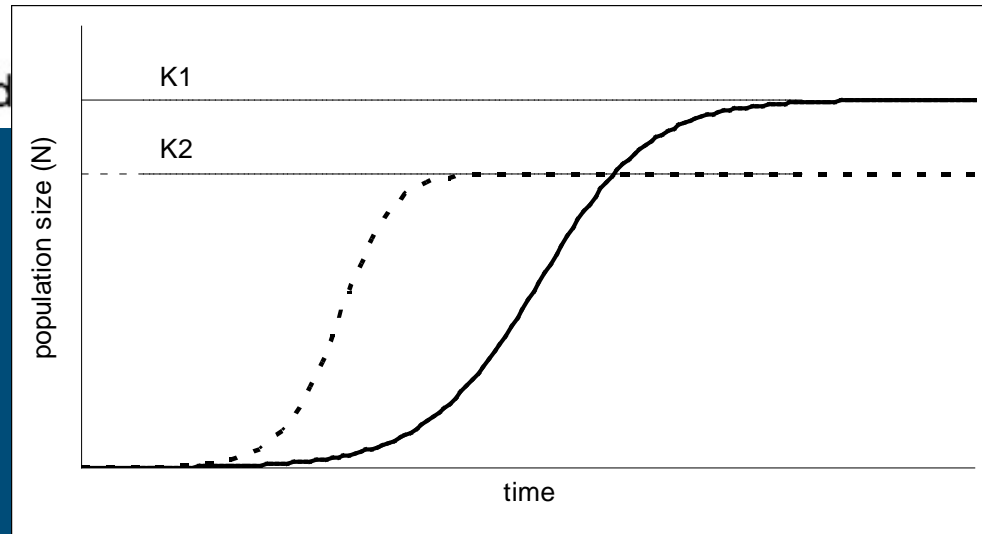
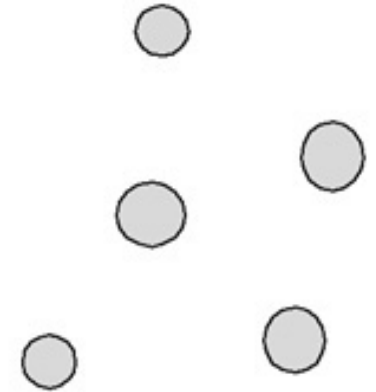
Demography in equilibrium model



(a) No subdivision



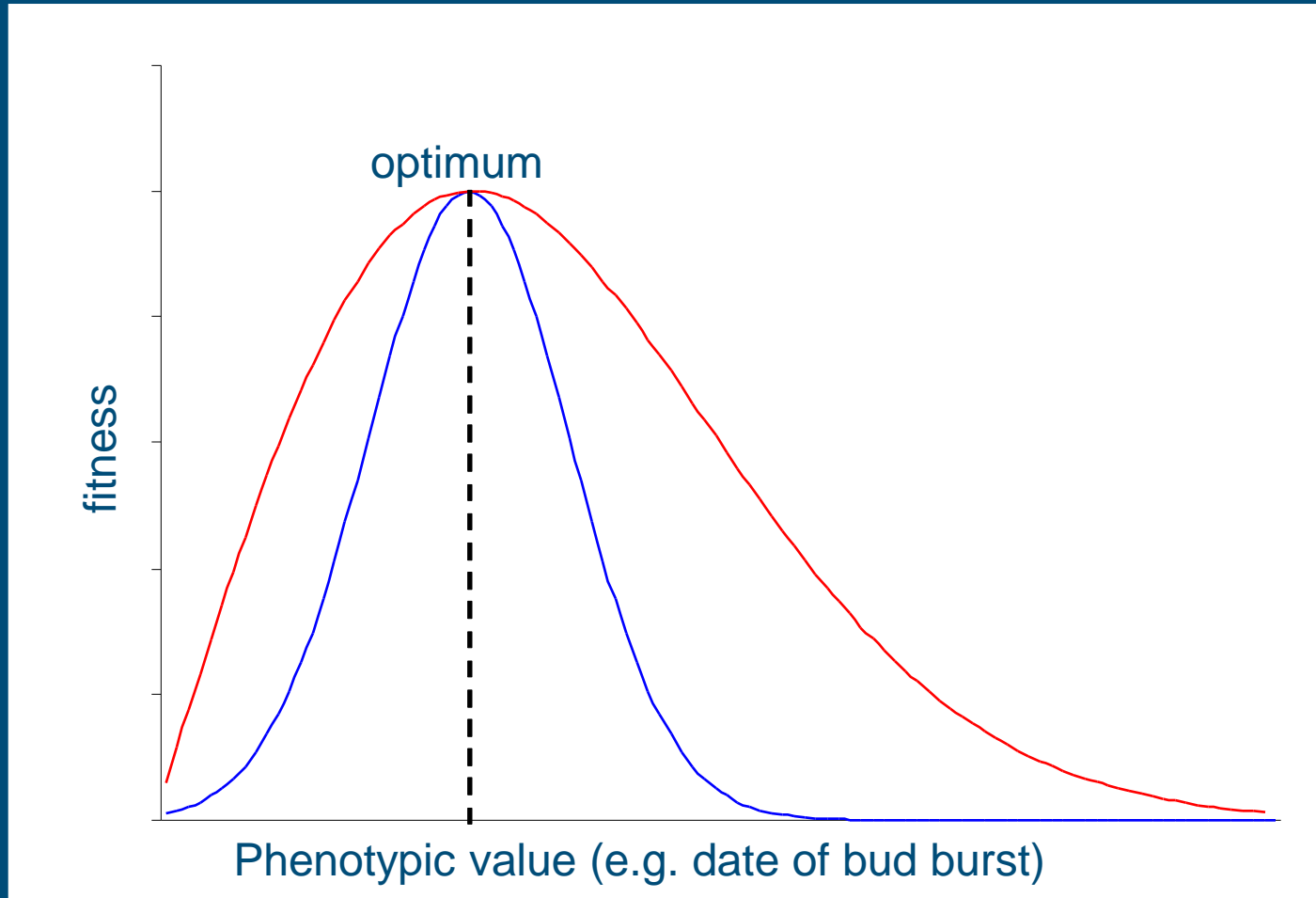
(b) Intermed



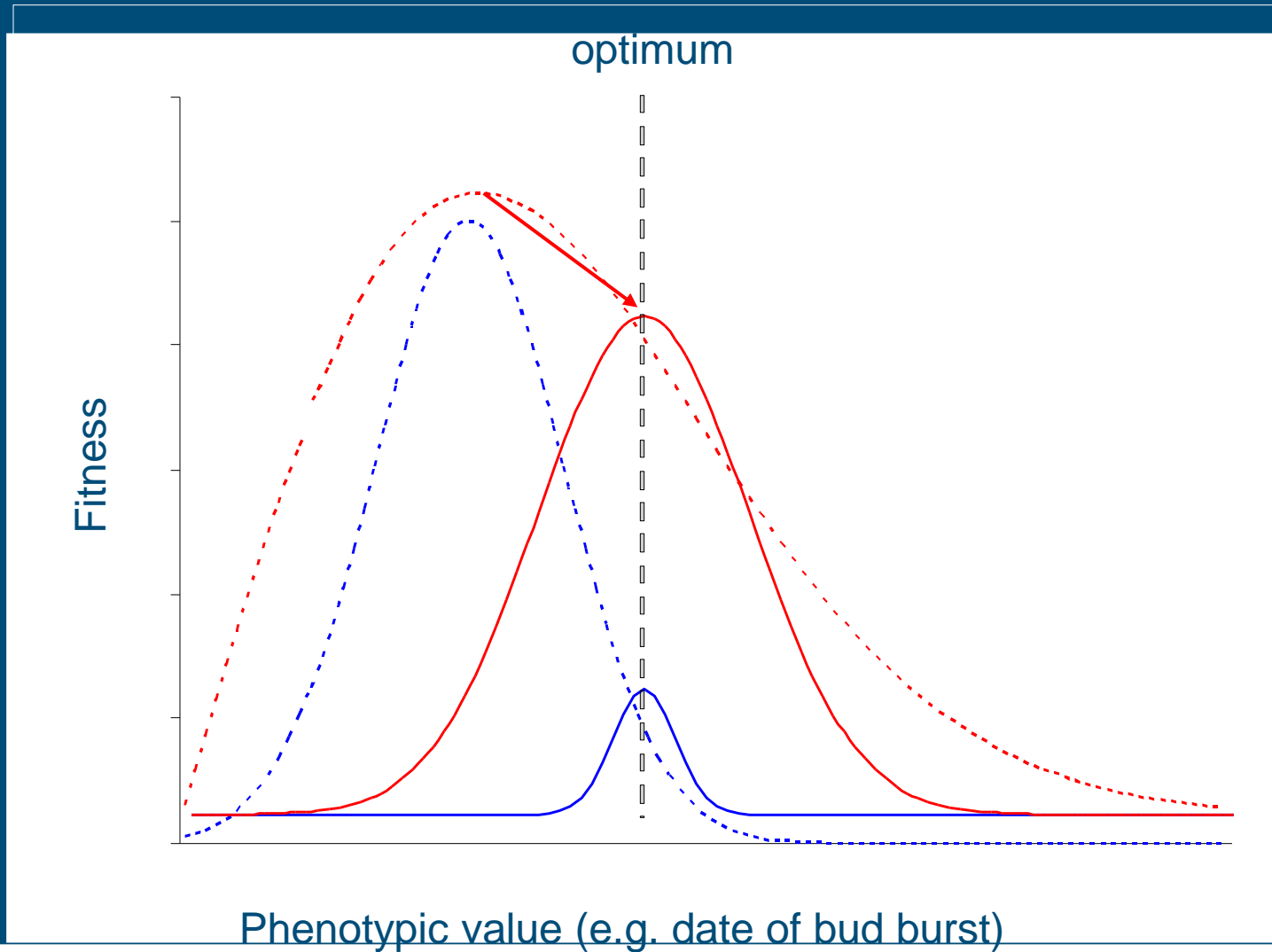
$$N_{t+1} = N_t + r \cdot N \cdot \left(\frac{K - N}{K} \right)$$

Classical population-genetic models – current situation

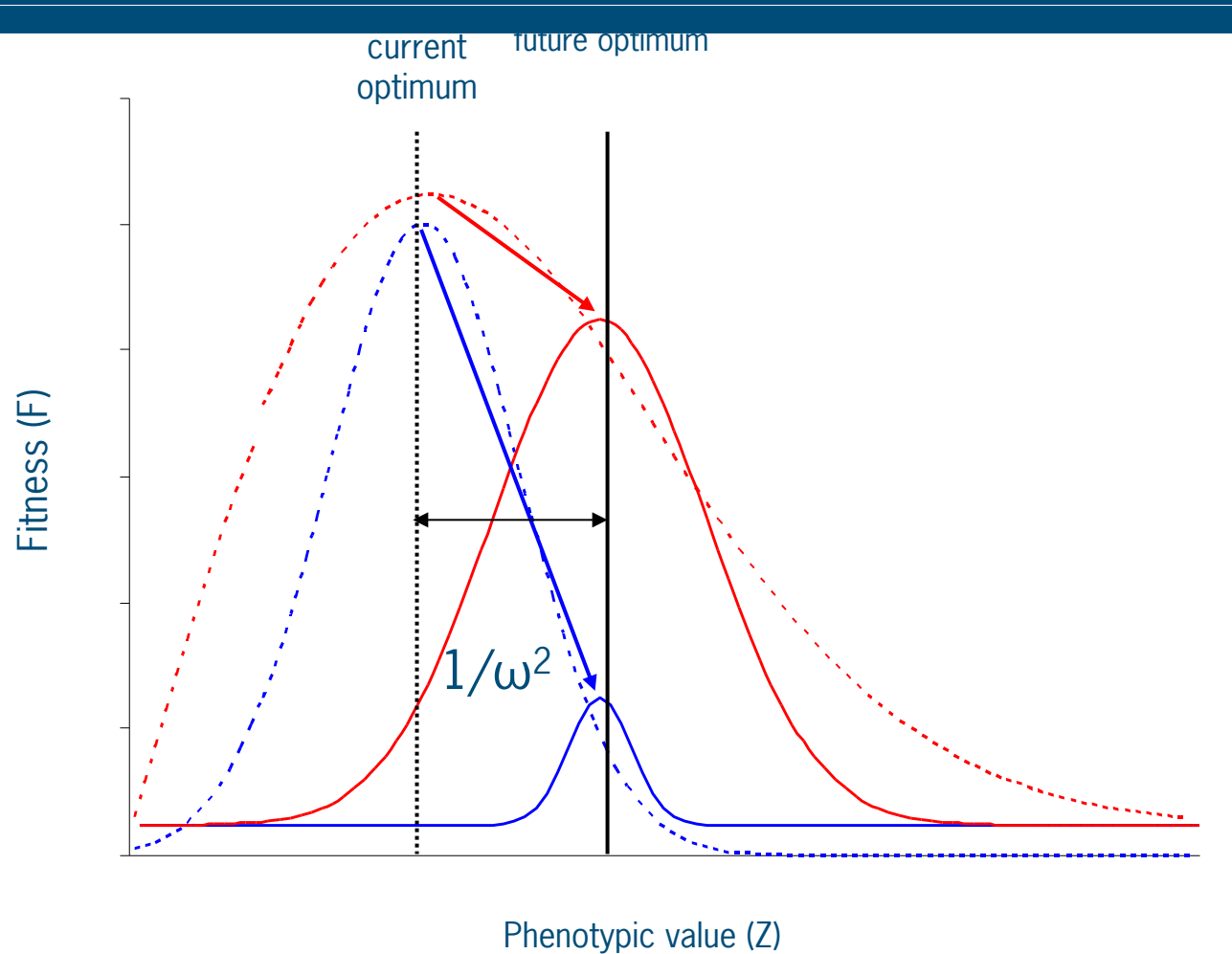
- 2 populations



Classical population-genetic models: future situation



Genetics in equilibrium model



$$F(Z) = \exp\left(-\frac{(Z - Z_{opt})^2}{2\omega^2}\right)$$

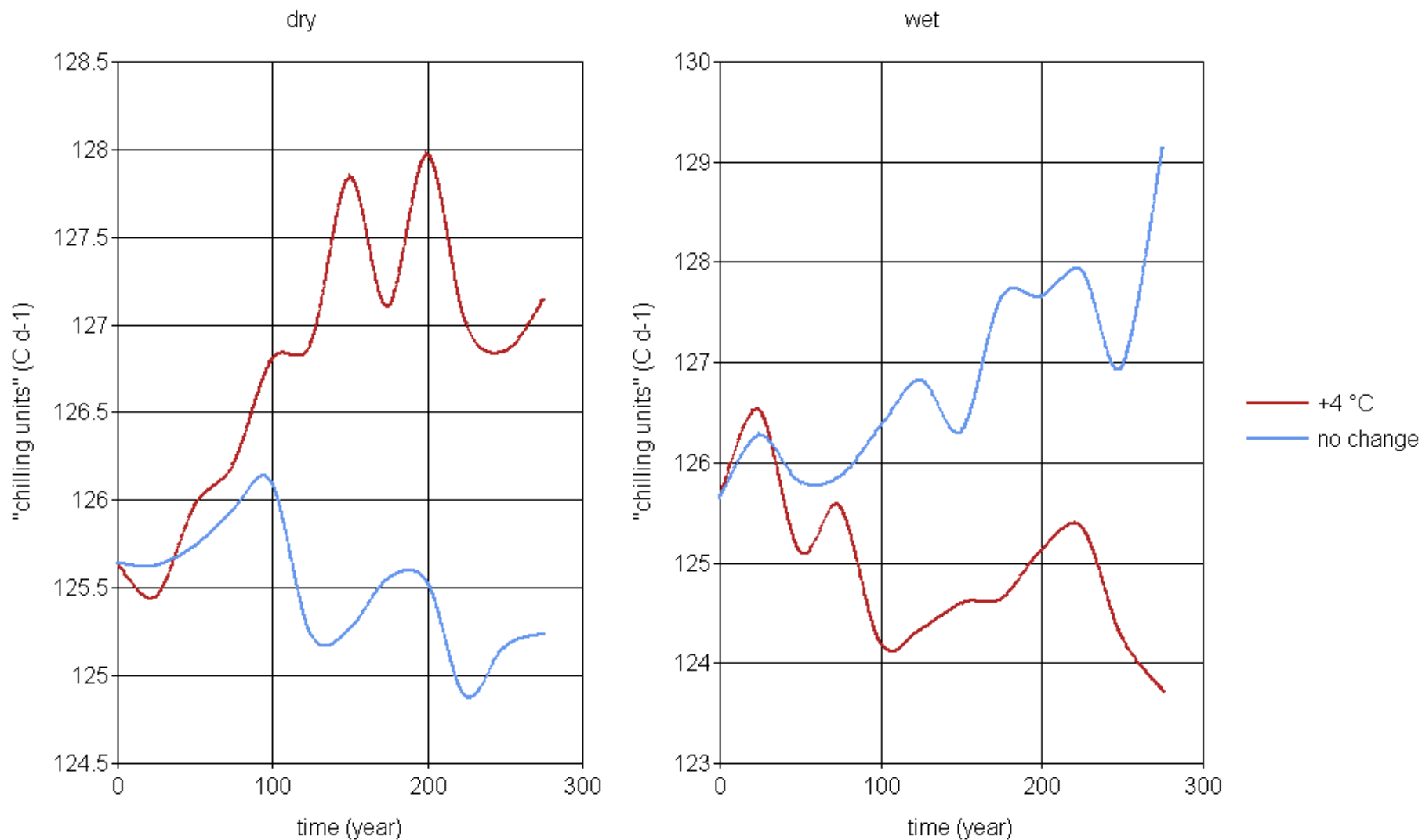
Non-equilibrium approach: individually-based genetic modeling



Non-equilibrium or individualistic approach

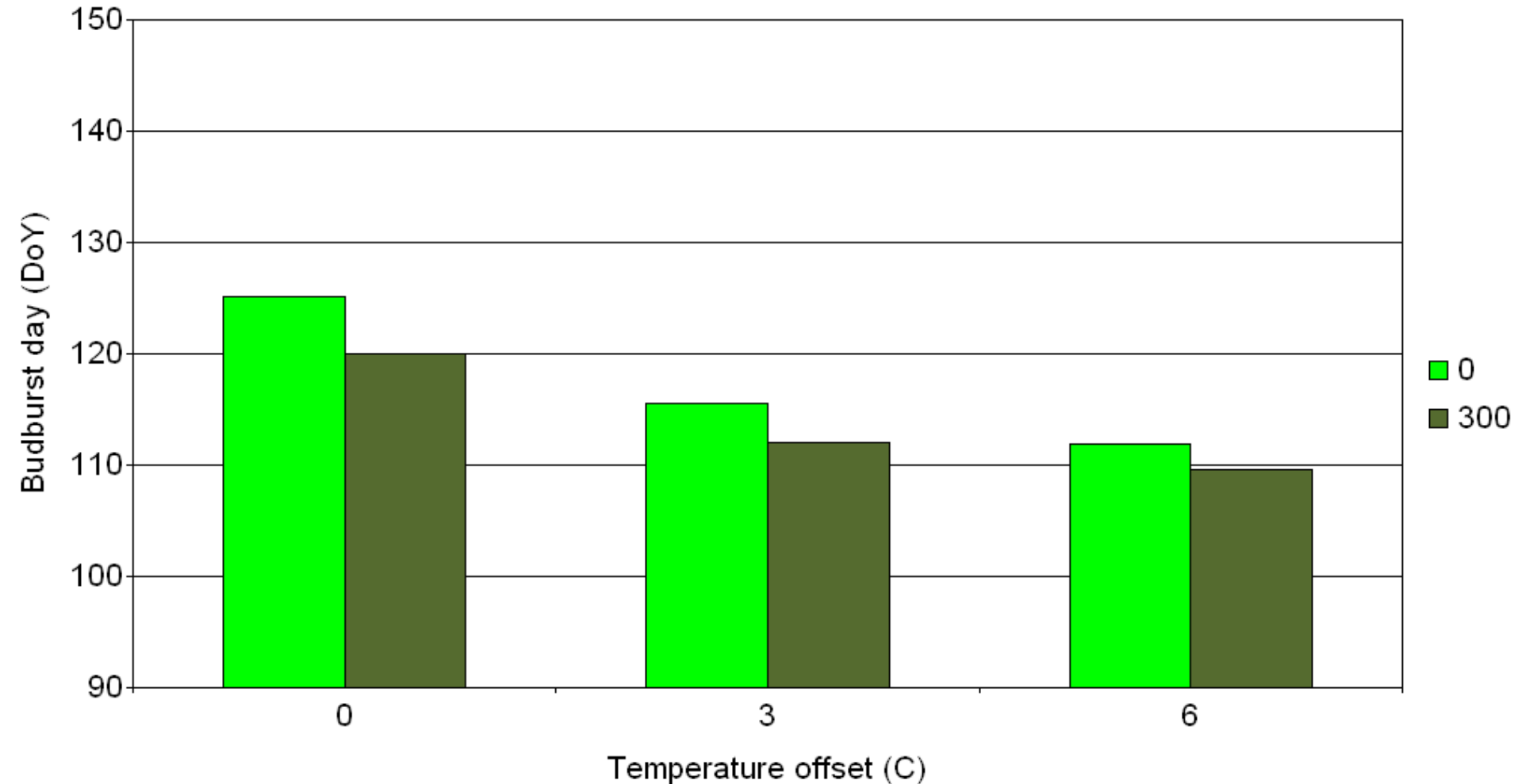
- Assumption:
 - Environment is non-stationary in space and time relative to the rate of adaptation
 - Population is always lagging behind changing biotic and abiotic conditions – both genetic and demography
 - => History does not provide knowledge on future “stable states”
 - => we have no information on future stable states
- Traits e.g.:
 - Budburst, growth, WUE, NPP, biomass, height
- Model parameters under study e.g:
 - Critical temperature thresholds, sensitivity of process to environmental driver
 - i.e. parameters that determine phenotypic plastic response but are assumed to be invariant with respect to environmental conditions
- Model analyses e.g.:
 - Determine processes and traits that are most under selection
 - Study change in phenotypic plasticity in (future) environmental conditions and assess role of spatial genetic structure, gene flow etc.

E.g. 1. Evolution of critical state of chilling (S_{chl}^*)

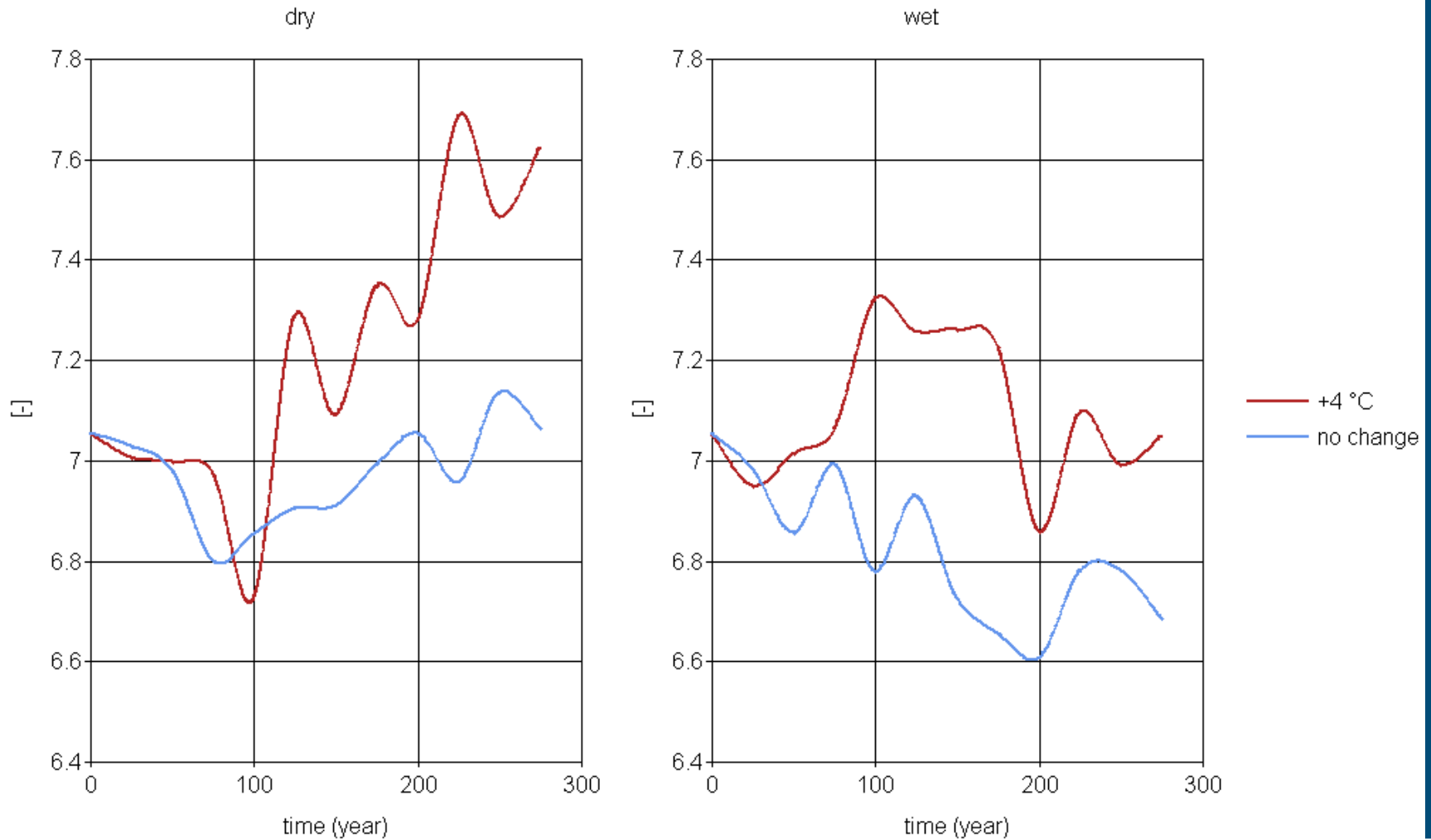


Consequence of change in S_{chl}^* on phenotypic plastic response of bud burst to temperature

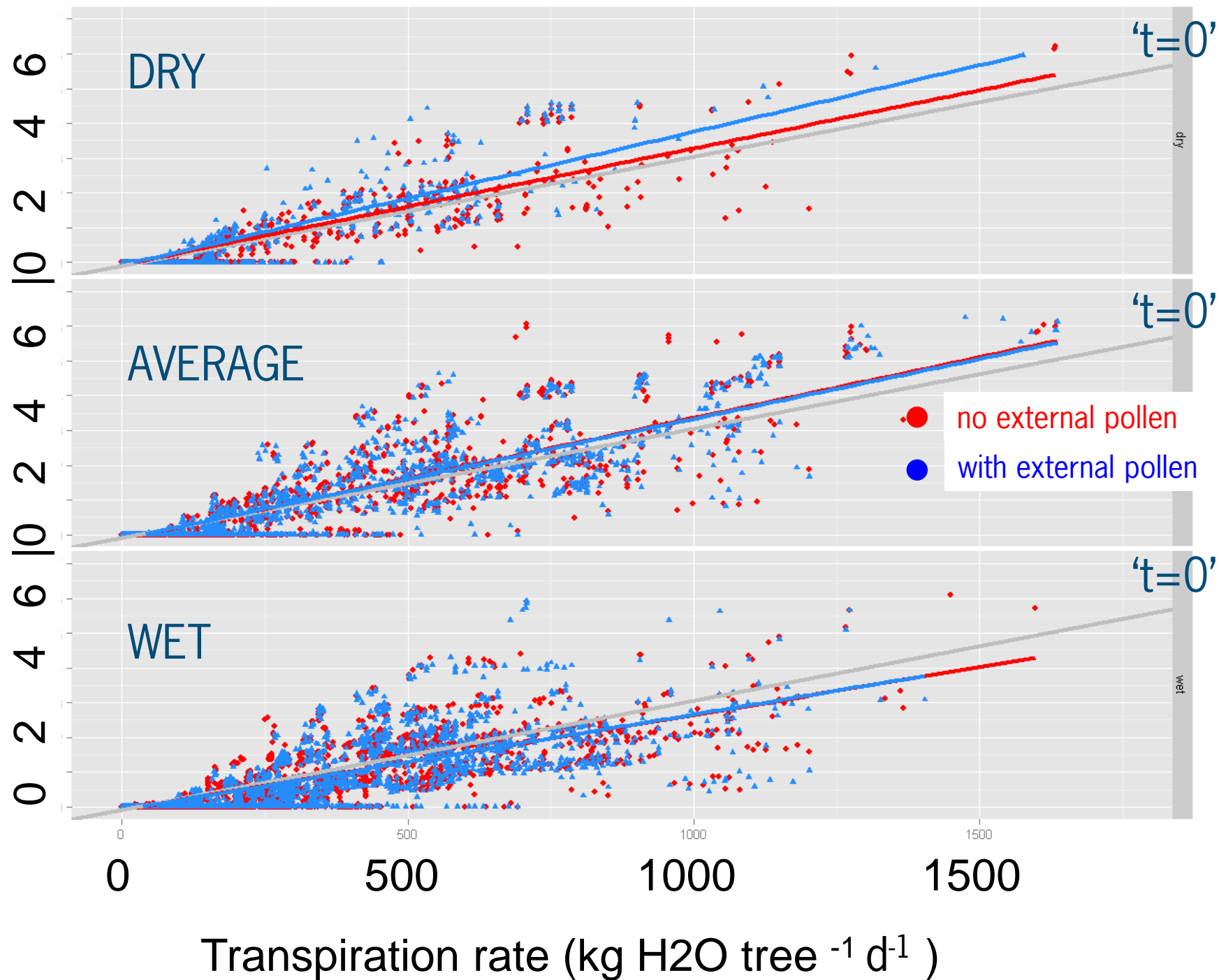
Phenotypic plastic response of bud burst to temperature at $t=0\text{yr}$ + adaptive response at $t=300\text{yr}$



E.g. 2: Evolution of sensitivity of stomatal conductance to soil water availability



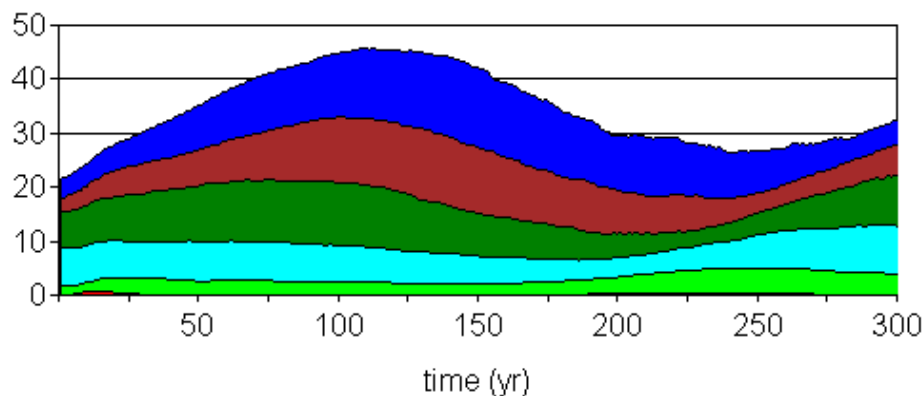
NPP (g C tree⁻¹ d⁻¹)



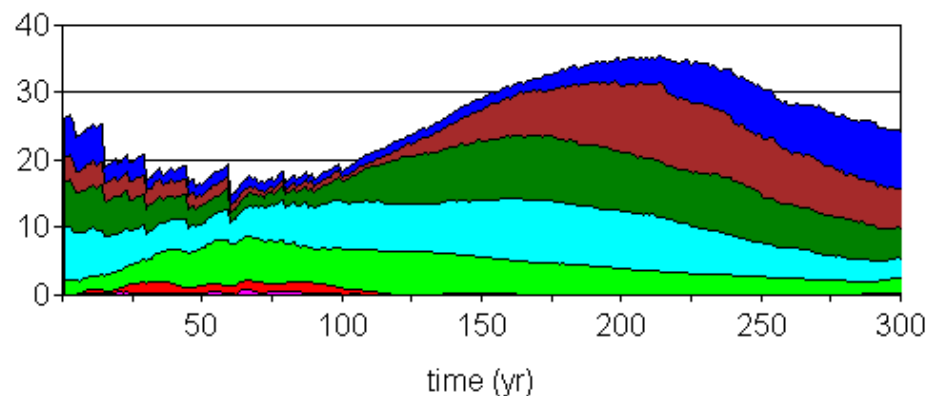
Example output ForGEM - Basal area

Basal area per Dbh-class (m² ha⁻¹)

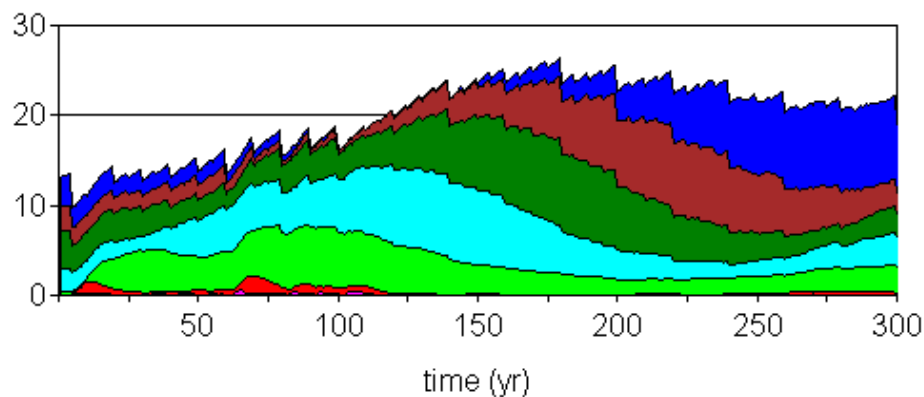
Fagus sylvatica - 1. No Management



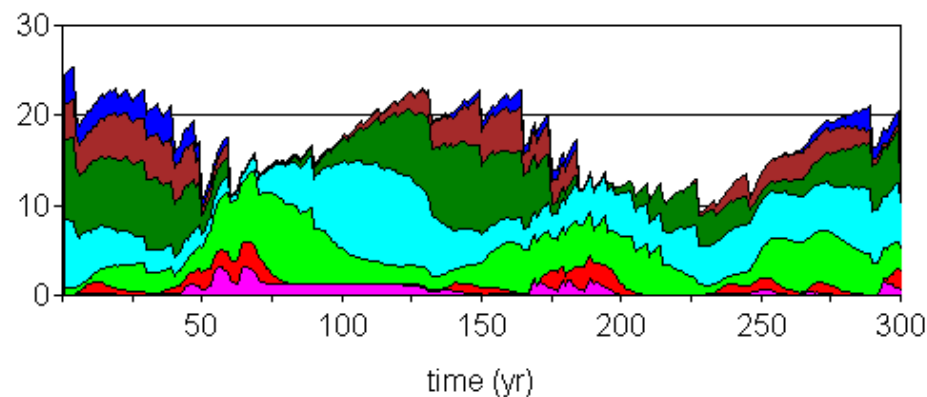
Fagus sylvatica - 2. Nature oriented



Fagus sylvatica - 3. Group selection



Fagus sylvatica - 4. Sheltercut



■ Average of Dbh90-500 ■ Average of Dbh70-90 ■ Average of Dbh50-70 ■ Average of Dbh30-50
■ Average of Dbh10-30 ■ Average of Dbh5-10 ■ Average of Dbh0-5

Example output ForGEM: Genetic diversity

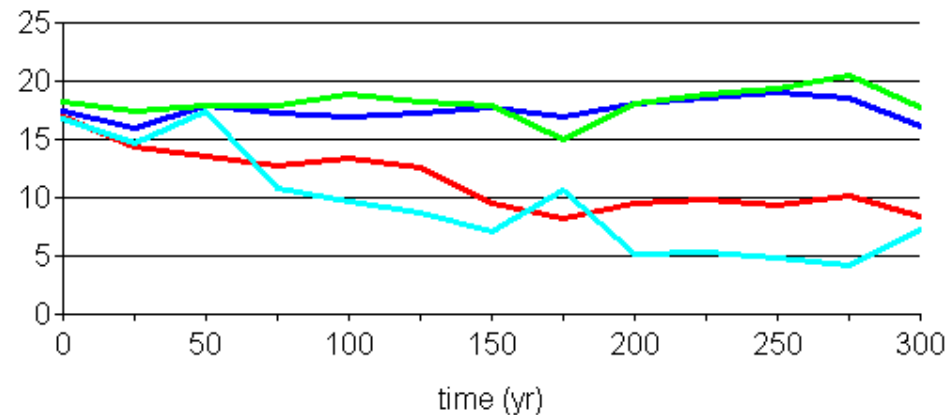
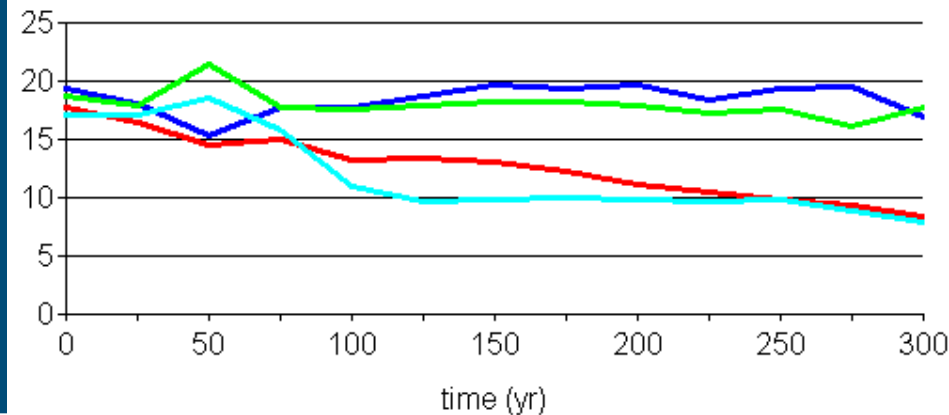
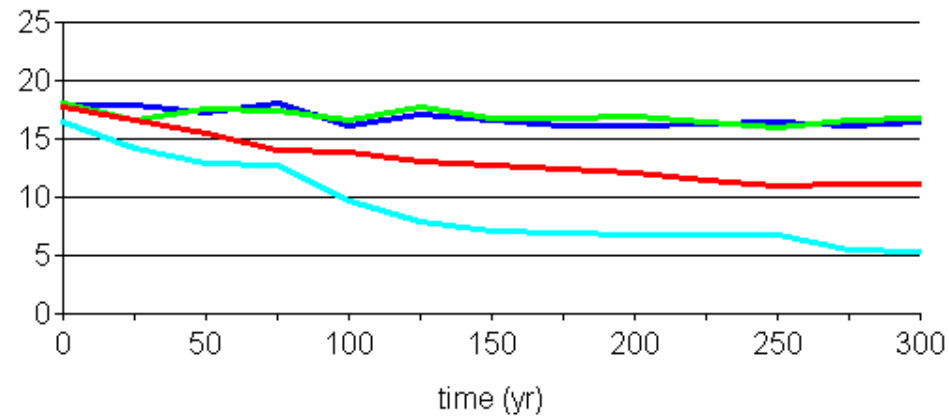
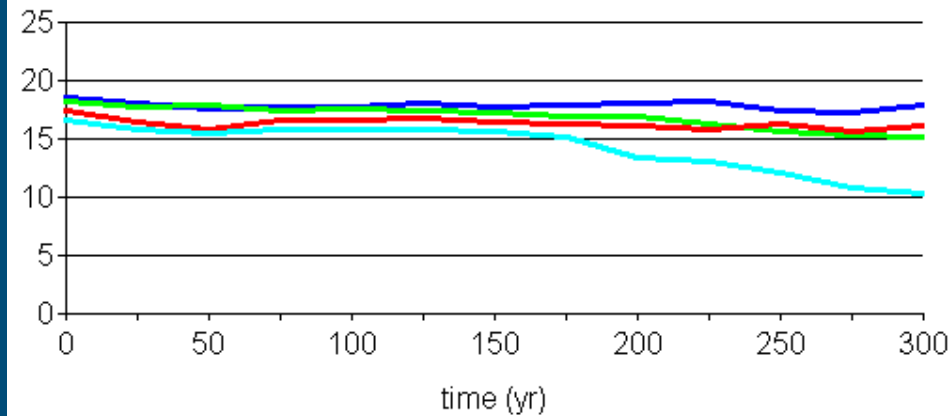
Genetic diversity

1. No Management

2. Nature oriented

3. Group selection

4. Sheltercut



— 1. Neutral trait — 2. Budburst day — 3. Spiral grain — 4. C7Hgh


Pros and cons of eq. and non-eq. genetic modeling

■ Eq.:

- Generic, suitable for analysis of past, long-term evolutionary processes
- Abstract traits related to whole tree fitness function
- Not suitable for short-term future assessment because equilibrium states and selection pressure are input to the model

■ Non-eq.:

- Realistic, suitable for prediction at short-term, also for future equilibriums
- Traits that have trade-off in resource use and fitness, that results in phenotypic plastic responses (morphological / physiological)
- Not suitable for long-term (>10s generations) evolutionary processes



Conclusions on adding a genetic model to existing forest model

- Don't add an equilibrium genetic model to a process-based forest models
- Don't add phenotypic traits with partial fitness effect in equilibrium model