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Local adaptation to climatic gradients: application of a Physio-Demo-Genetics simulation model to **European Beech populations** Capsis4

Écologie des Forêts Méditerranéennes **INRA** Avignon

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The physio-demo-genetic model



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Predicting the response of trees, the keystone elements of forests, to ongoing climate change is a critical issue.

Forests are the most important terrestrial ecosystems, providing 75% of the terrestrial biomass and several ecosystem services. Climate change has already taken its toll on forests and the future of forests is unforeseen. Trees are key drivers of ecosystem processes and stability in forests, directly sustaining diverse plant and animal communities, and mediating nutrient and water ecological cycles Over the past two decades, extensive forest tree mortality triggered by droughts and extreme climatic conditions has been documented all around the world. Nonetheless, tree populations response to climate is still not sufficiently well understood to be able to make reliable predictions for the future. Besides migration, Forests response to climate will be determined by a combination of factors such as phenotypic plasticity and adaptive capacity that are largely unknown for many tree species.

We propose a new Physiological-Demographic-Genetic (PDG) coupling:

1) an eco-physiological and physical module (CASTANEA) simulating carbohydrate and water fluxes at tree-level based on daily climate observations (real or simulated).

2) a population demographic module

converting carbohydrate reserves into demographic rates for adult tree (growth, mortality and seed production) and simulating ecological processes across the life cycle (including seed and pollen dispersal, germination success, density-dependent mortality of seedlings).



Here we develop a new simulation approach to investigate the evolutionary potential of forest trees to adapt to ongoing climate change. We build an individual based model that accounts explicitly for the effect of individual genetic variability and climate on the variation of tree physiological traits. Selection is acting on functional traits involved in response to climate, such as the timing of budburst (TBB). We illustrate how this tool can be used to investigate the rate of evolutionary change in tree populations evolving for few generations along an environmental gradient.



3) a quantitative genetic module (GENETICS) relating the genotype at quantitative trait loci (QTLs) to the phenotype of one or more functional traits, accounting for environmental effect. Individual fitness is an output of the model, calculated as the life time reproductive success resulting from a combination of functional traits.

Here we consider only the Timing of Budburst (TBB) = f (Sum of degree days, Temperature)





