

WHERE DID ARCTIC-ALPINE MOSSES SURVIVE IN A FROZEN EUROPE?

Insights from a multispecies
coalescent analysis

Alice Ledent, Aurélie Désamoré, Benjamin Laenen, Jairo Patiño, Stuart F. McDaniel,
Patrick Mardulyn and Alain Vanderpoorten

Introduction

Impact of global warming on species distribution ?

→ Study of past climate changes (Petit, R.J. *et al.*, 2005)



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What does explain the actual species distribution ?

→ Quaternary glacial periods (Hewitt, G.M. 1996, 1999, 2000)

- Last Glacial Maximum (LGM: 26,000 – 19,000 years BP) = the most virulent

Fundamental biogeographic hypothesis in Europe

- Impact of LGM on species distribution
- Southern *refugia* hypothesis
- Temperate species
- Small mammals (Hewitt, G. *Nature*, 2000)
- Woody plants (Petit, R.J. *Science*, 2003)

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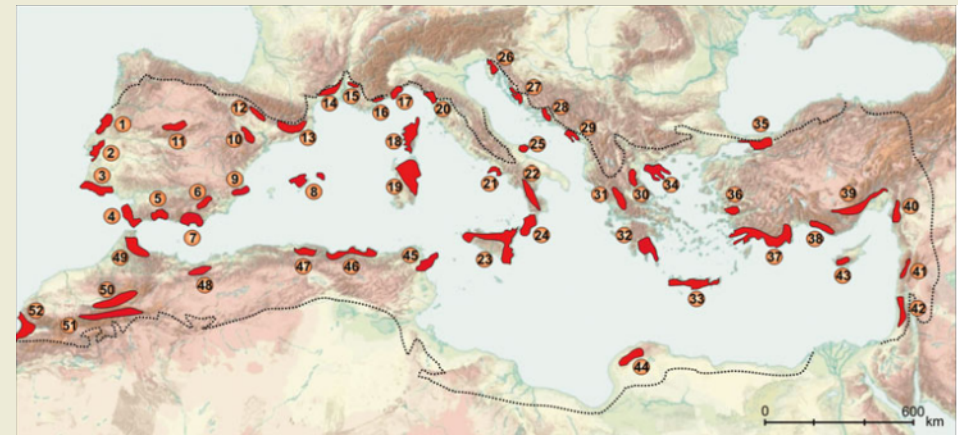
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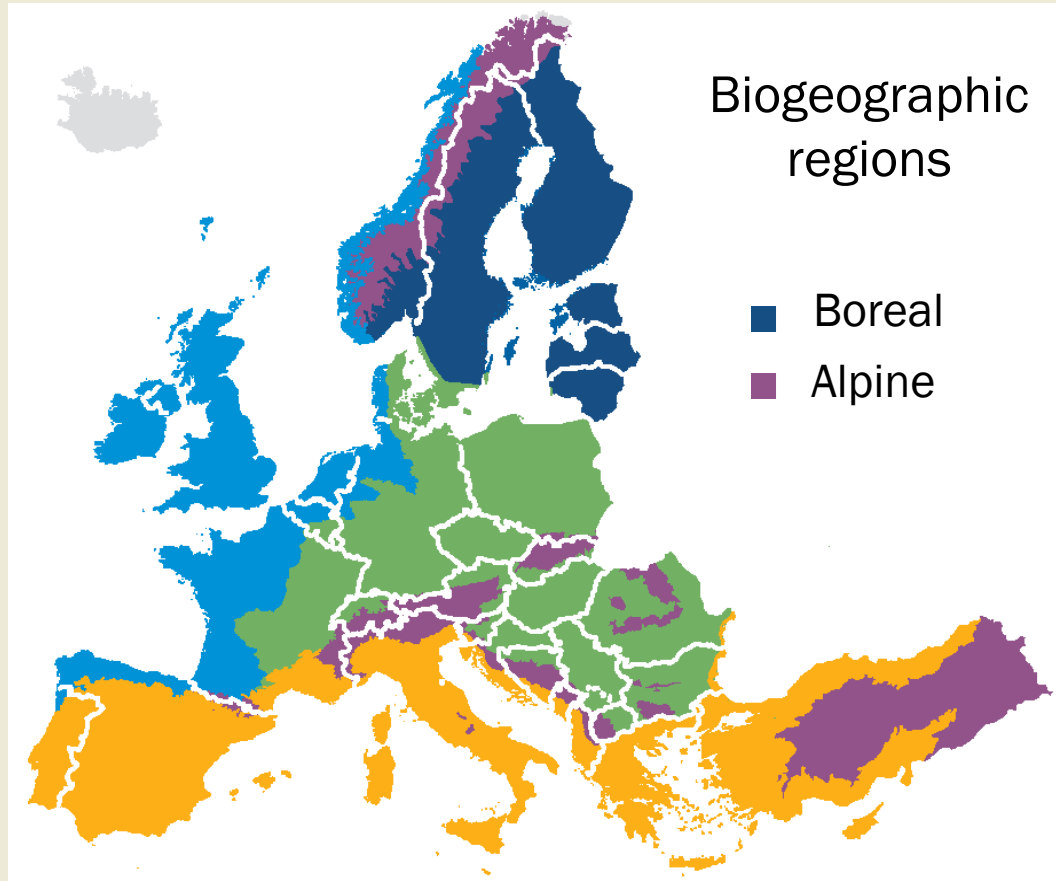
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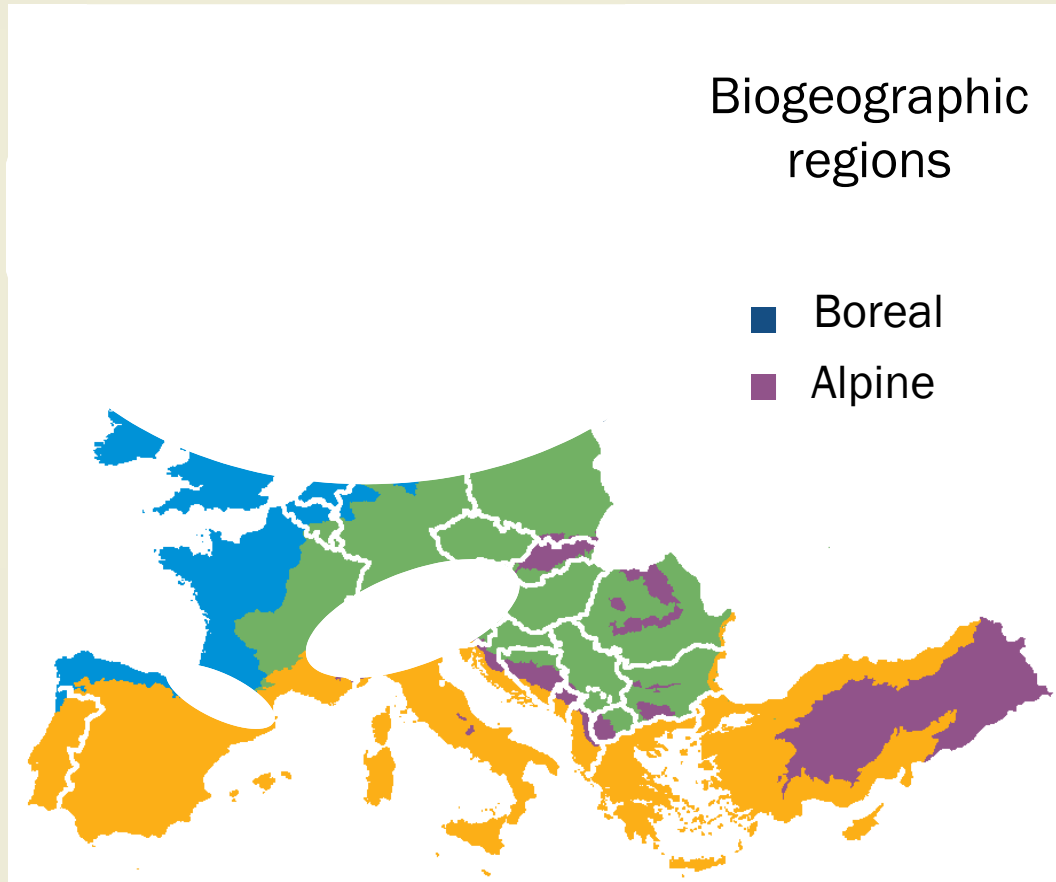
■ Arctic-Alpine distribution area



- Regions mainly covered by ice during the Pleistocene glacial periods
 - Disjunct distribution
 - *Arctic/Boreal = Fennoscandia*
 - *Alpine = Alps, Carpathians, Rhodopes,...*
 - Poorly studied but highly important
- Where did they survive during the glacial periods, especially the LGM?

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■ Arctic-Alpine distribution area



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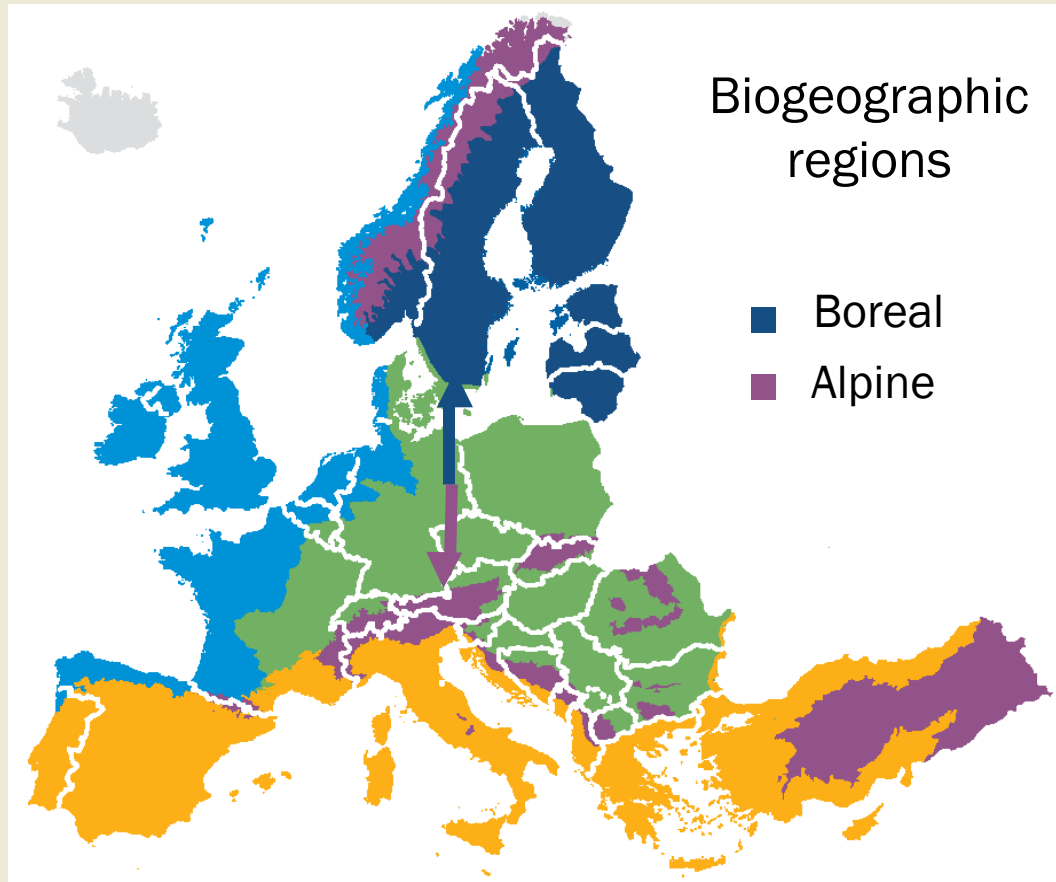
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Introduction

Major biogeographic hypotheses

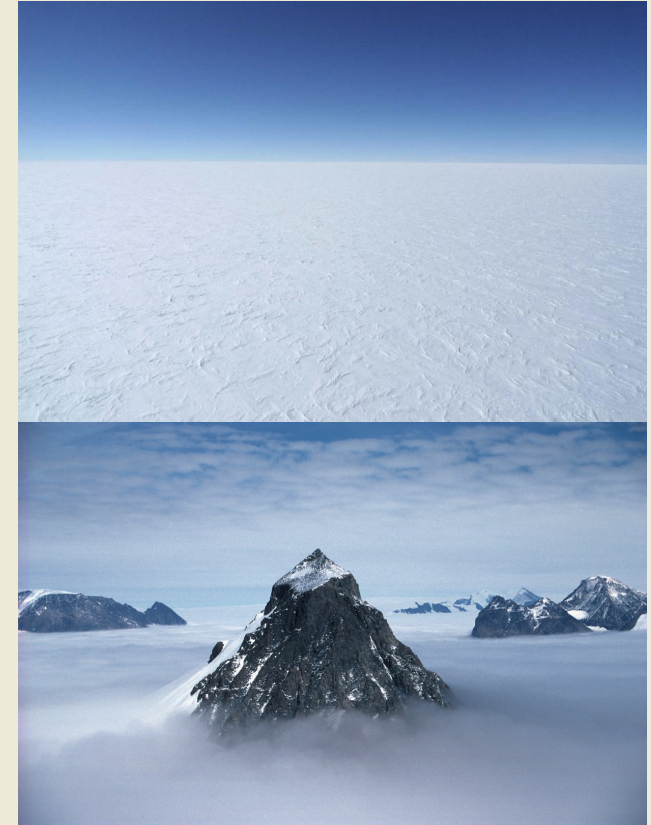
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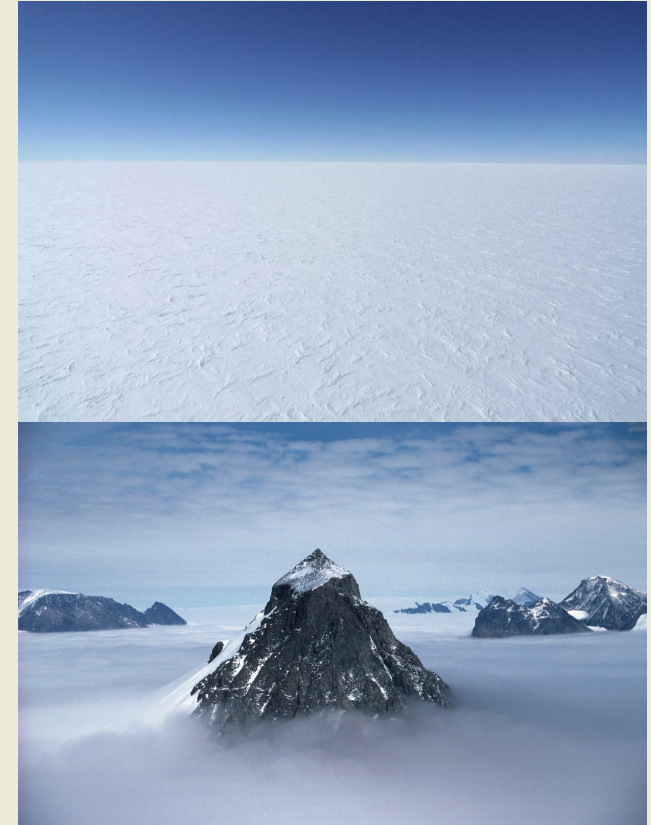
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Introduction

What about bryophytes?

- Dominant elements in Arctic-Alpine vegetation (Roads, E. 2014)
 - High cold tolerance (Furness, S.B. and Grime, J.P. 1982)
 - Ability to survive in ice and regenerate (Lafarge, C. 2013, Roads, E. 2014)
- Good candidate for the *Nunatak* hypothesis

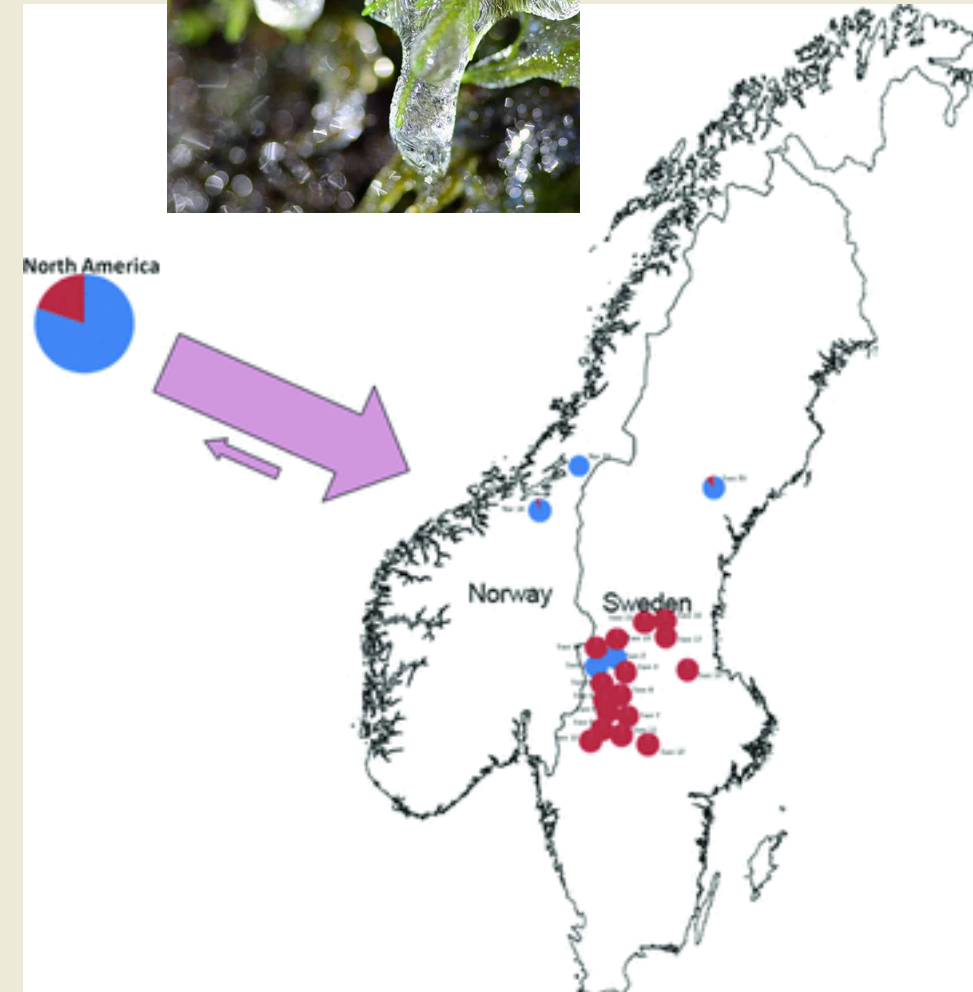
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Material and methods

I

Sampling and data analysis

- 3 species
 - *Amphidium lapponicum*
 - *Timmia austriaca*
 - *Timmia bavarica*
- Sampled across 5 populations
 - Fennoscandia ()
 - Iced Alps ()
 - Non iced Alps ()
 - Lowland ()
 - Out (not represented here)
- 3-4 chloroplastic and nuclear *loci*/sp.

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Material and methods

I

Sampling and data analysis



II

Approximate Bayesian Computation analysis (ABC)

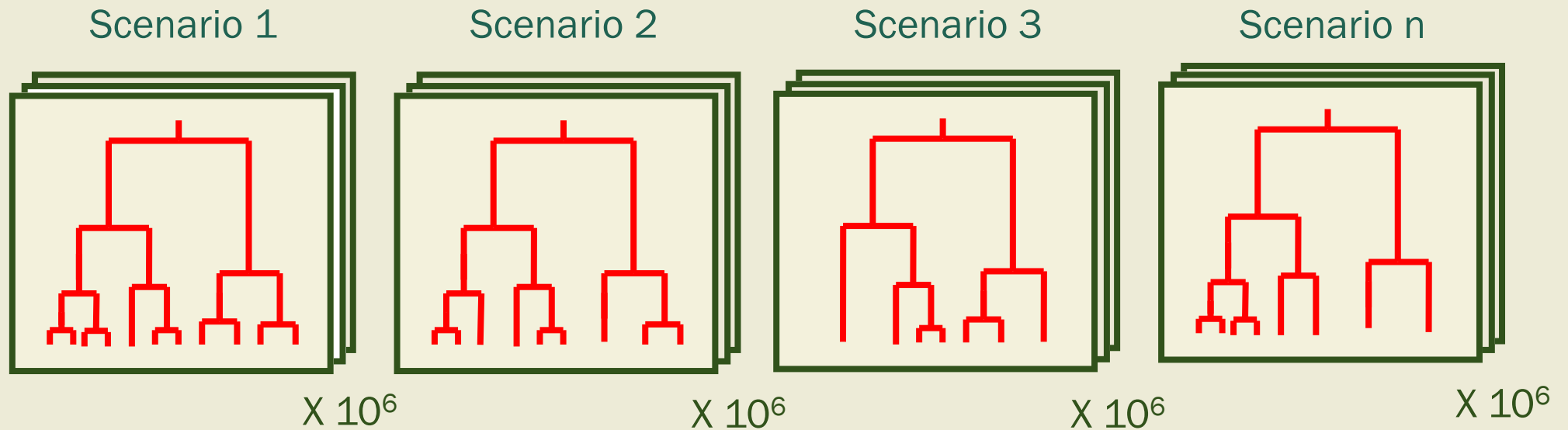


3 steps

Material and methods

1. Simulation of alleles genealogies

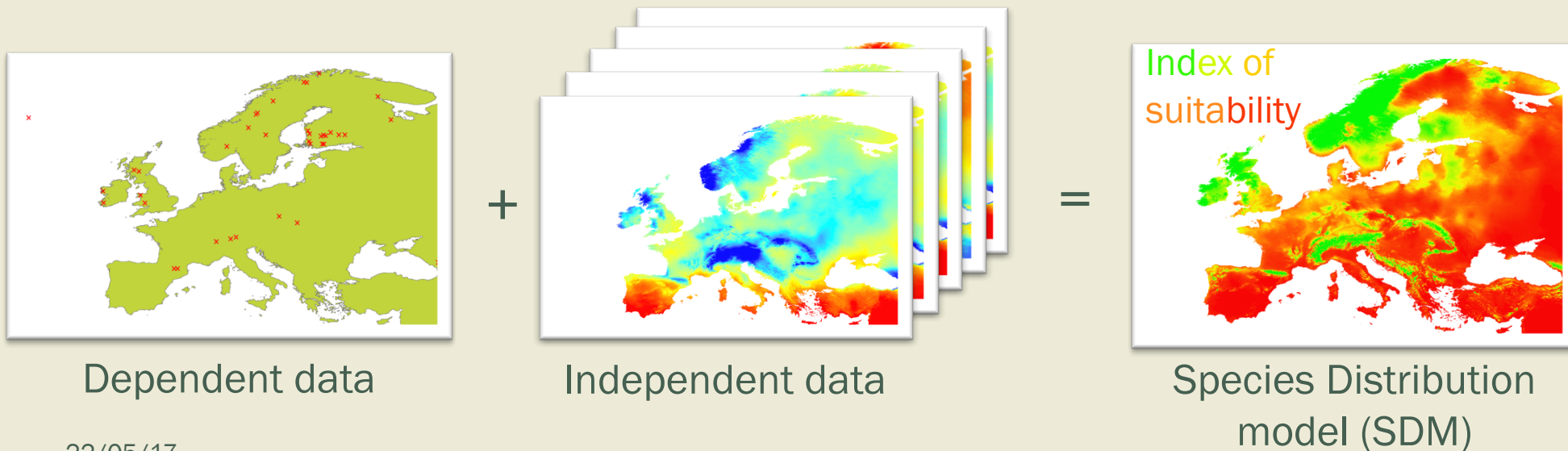
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- Under the constraint of different demographic scenarios
- Through definition of *prior* range of values of demographic parameters
 - *Migration rates*
 - *Effective population size*



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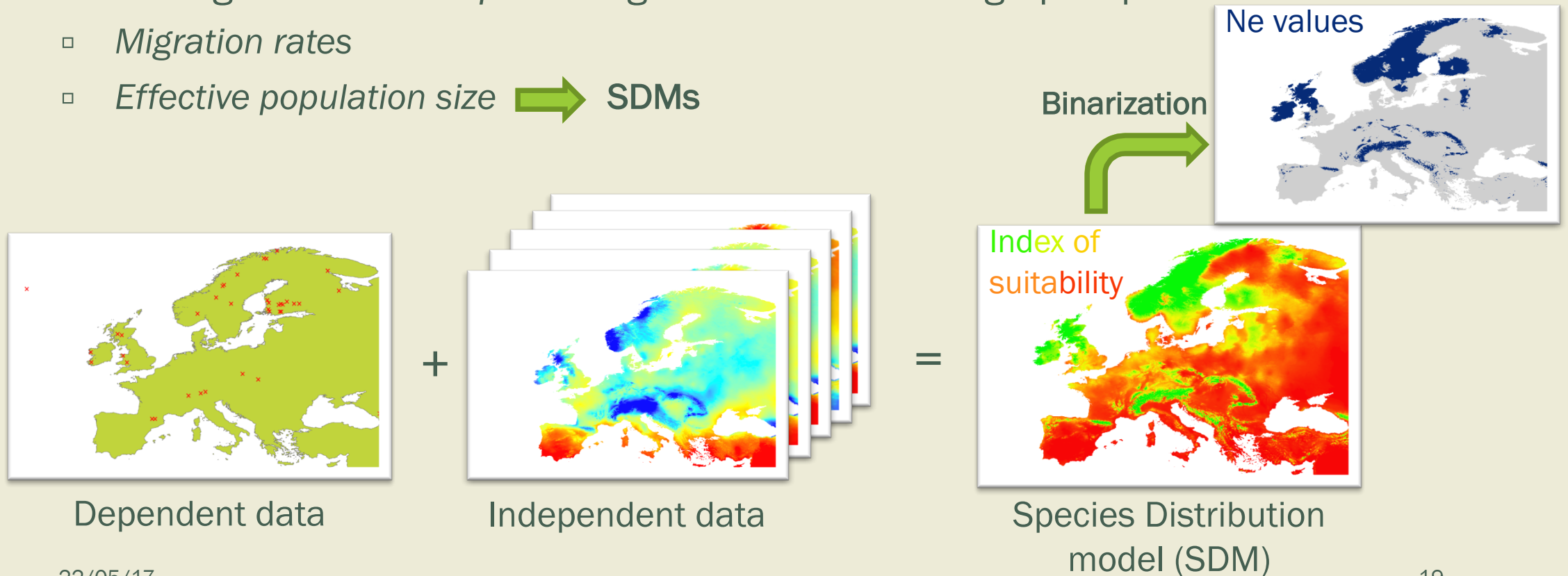
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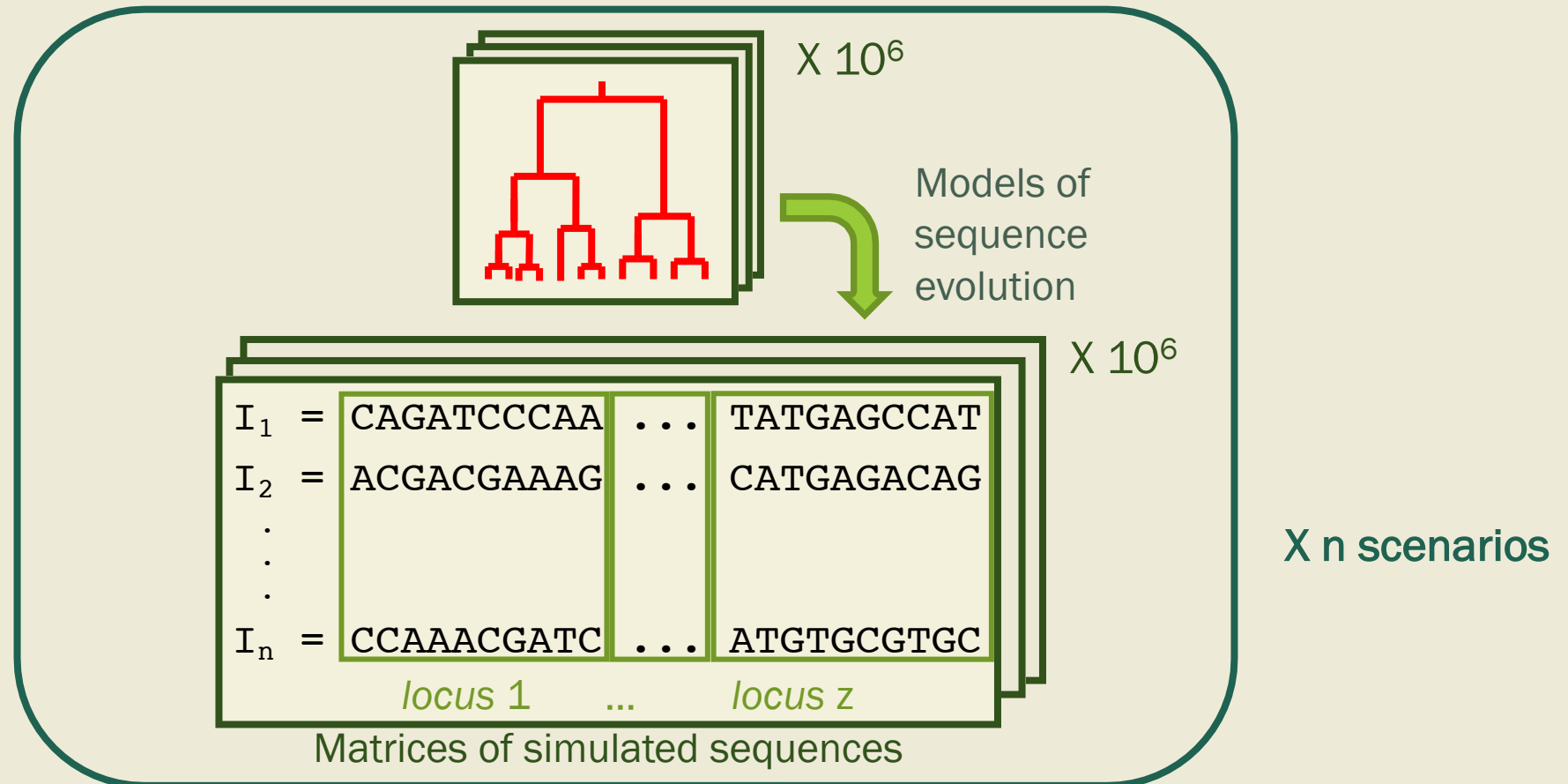
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 - *Effective population size* → SDMs



Material and methods

2. Matrices of sequences simulation

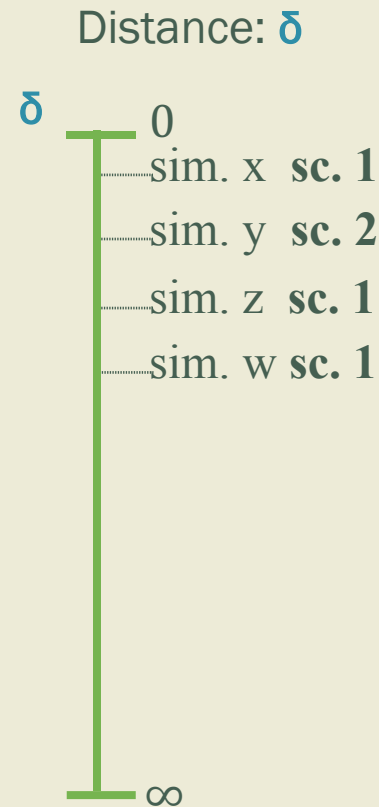
- Simulation of nucleotide matrices along each of the demographic genealogies using substitution models



Material and methods

3. Selection of the best-fit scenario

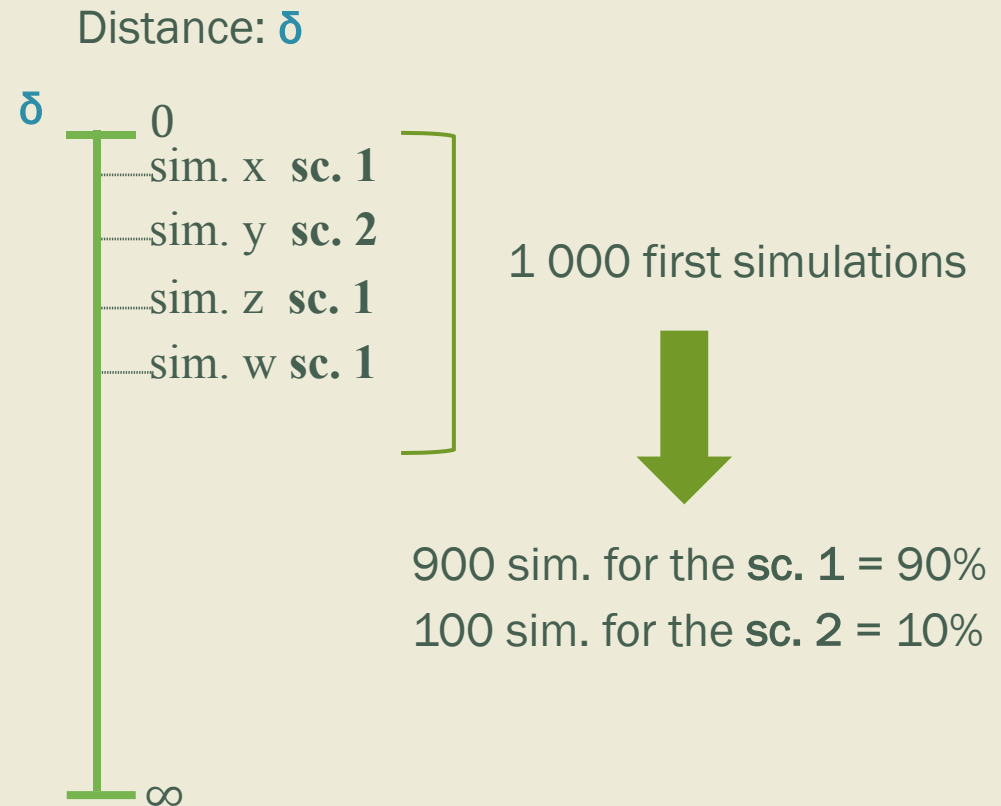
- Summary statistics: describe both observed and simulated datasets with descriptive statistics
- Euclidian distance: compute distance between each simulation and the observed dataset and rank simulations
- *Posterior probability*: determine, among the 1,000 first simulations, the proportion of simulations produced by each scenario
- *Best-fit scenario*: select the scenario with the highest *posterior probability*



Material and methods

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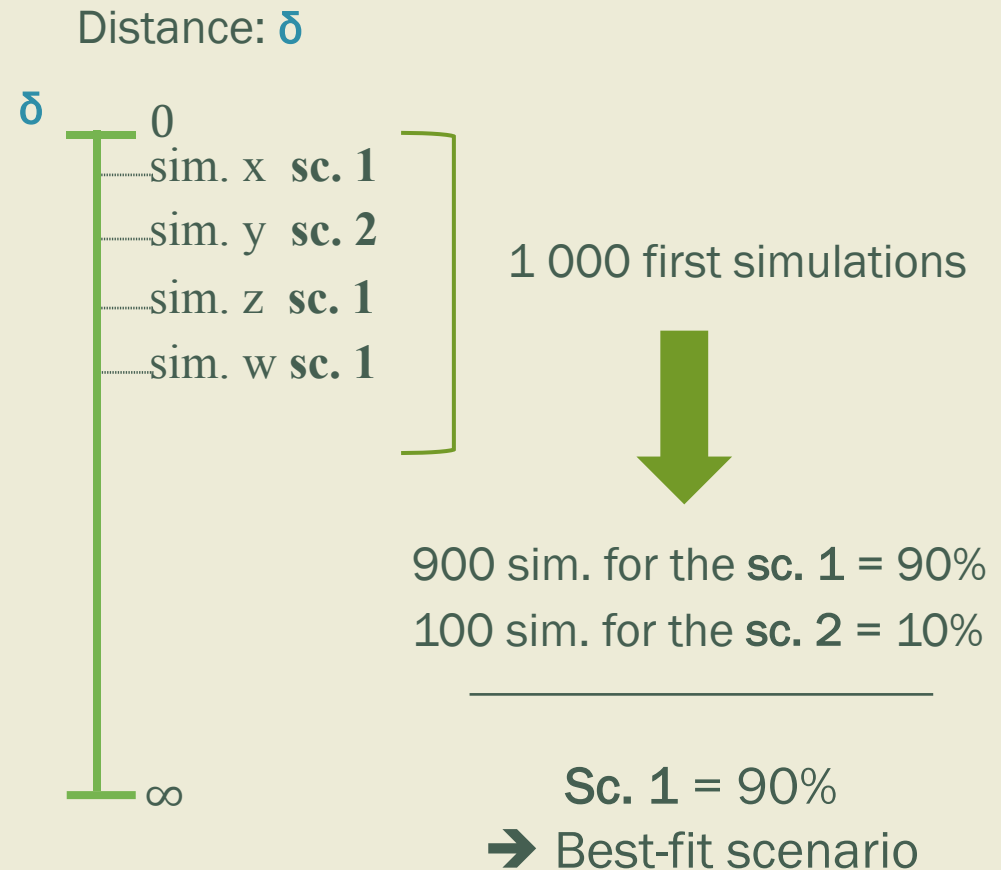
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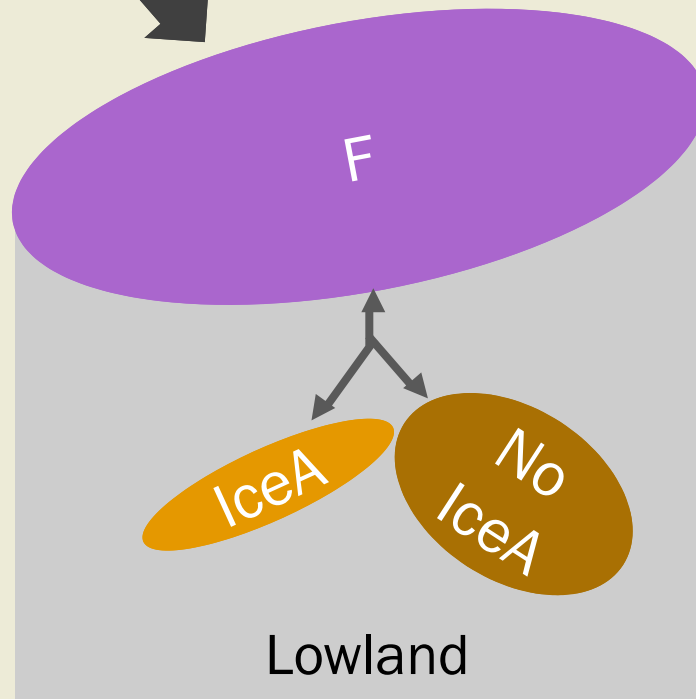
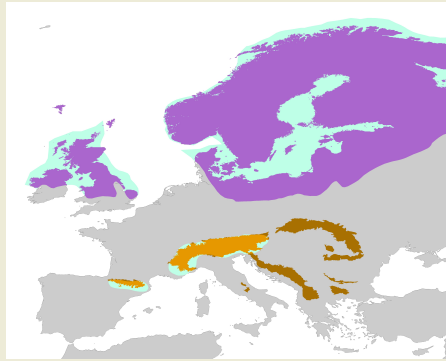
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Demographic scenarios




Effective population size

 = Empty

 = Colonization in progress

 = Full

Migrations = 

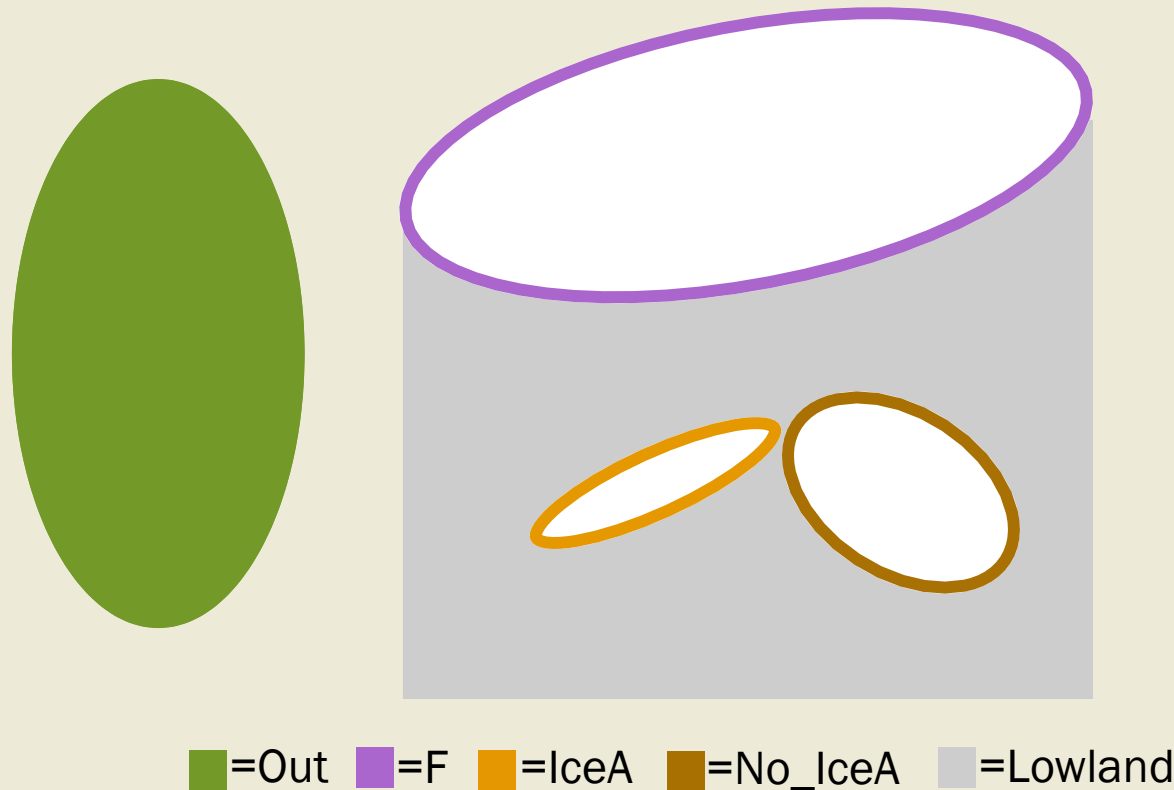
Periods

- LGM
- Onset
- Present

Demographic scenarios

Tabula rasa scenario

LGM



LGM

- No survival within the ice sheet
- Lowland areas suitable

Onset

- Recolonization from Lowland areas (outside the ice sheet)

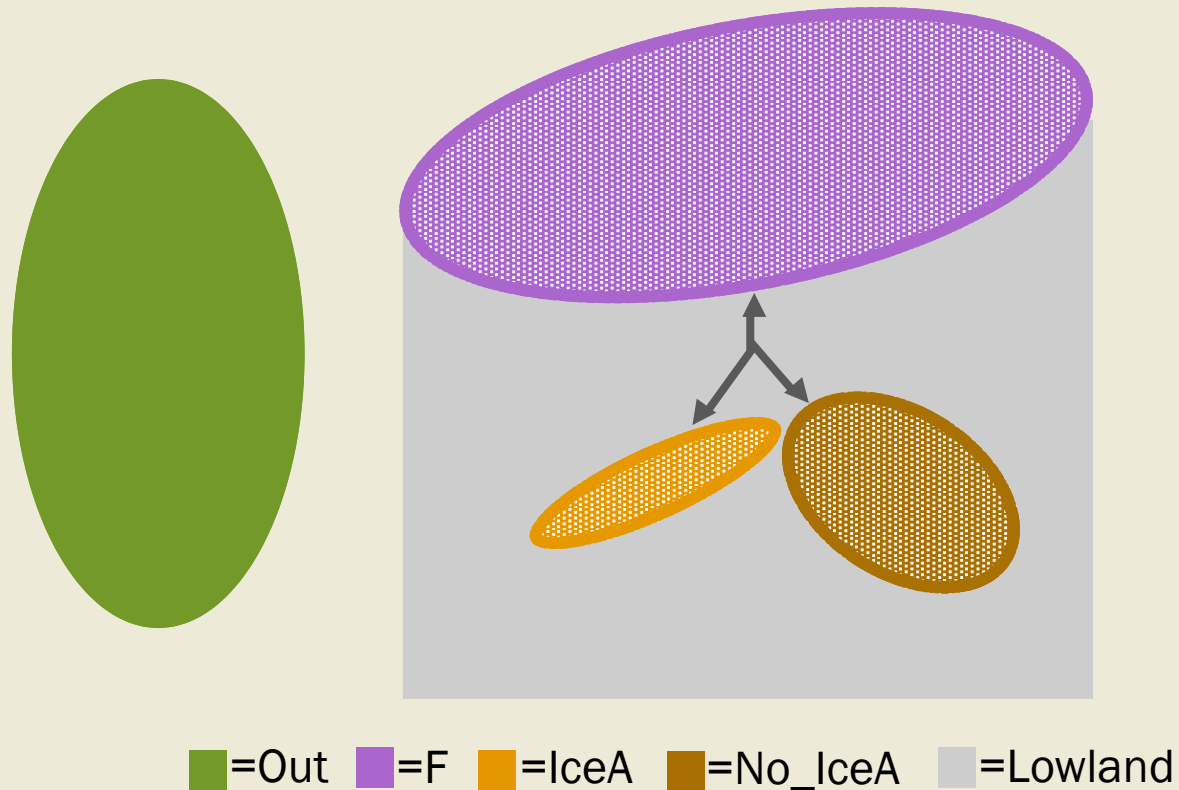
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- Lowland area no longer suitable
 - Too hot and dry
 - Too much competition

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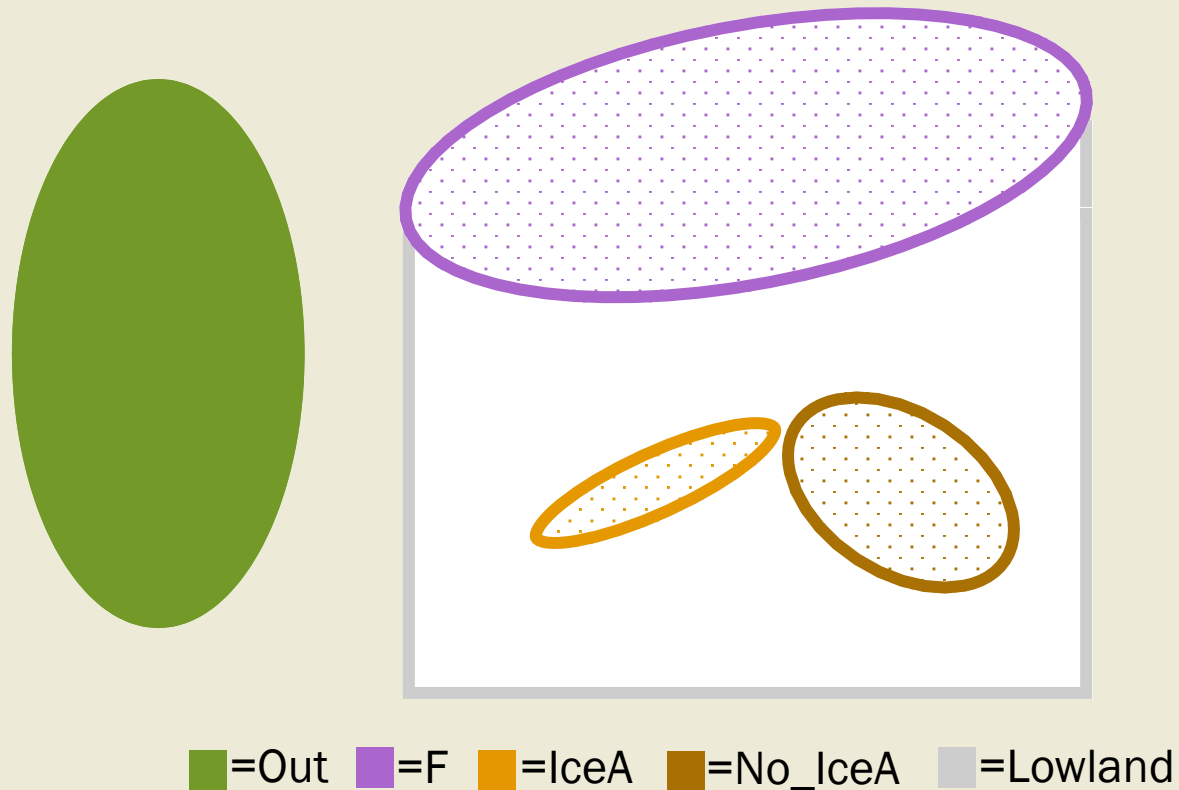
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Nunatak scenario

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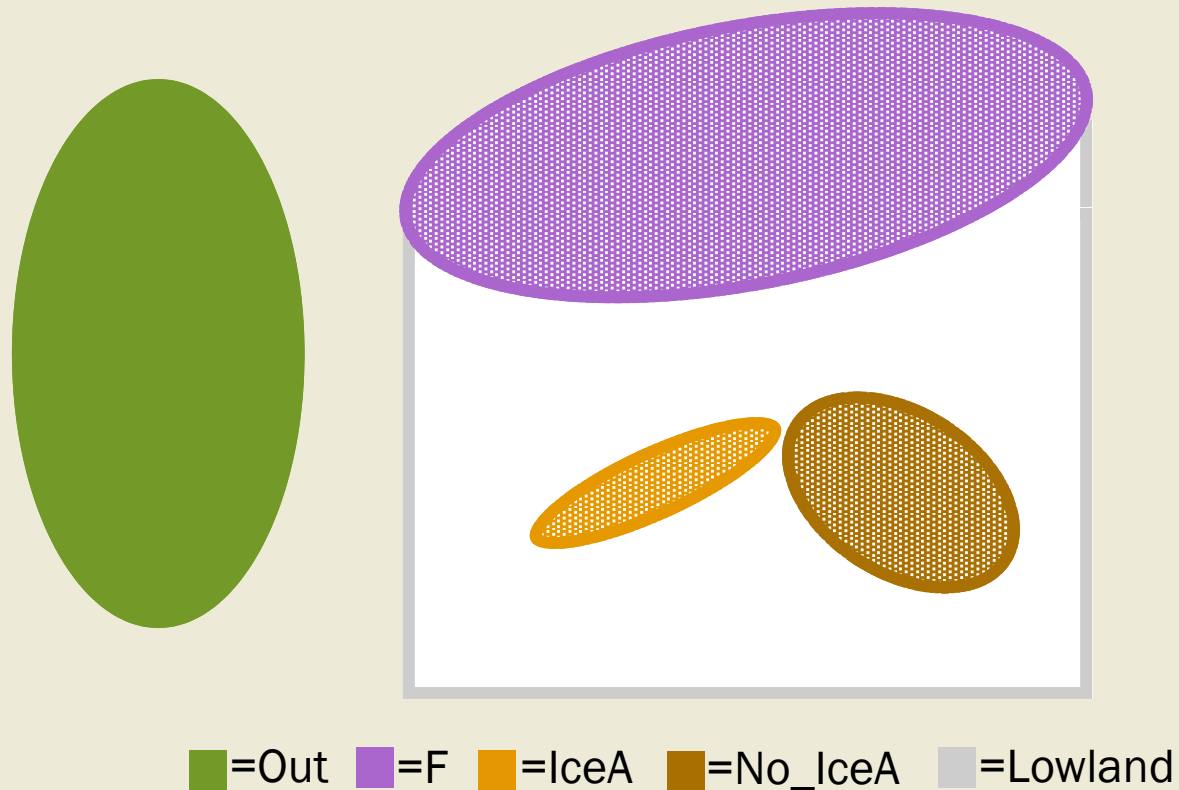
Onset

- Populations expansion from those refugia

Demographic scenarios

Nunatak scenario

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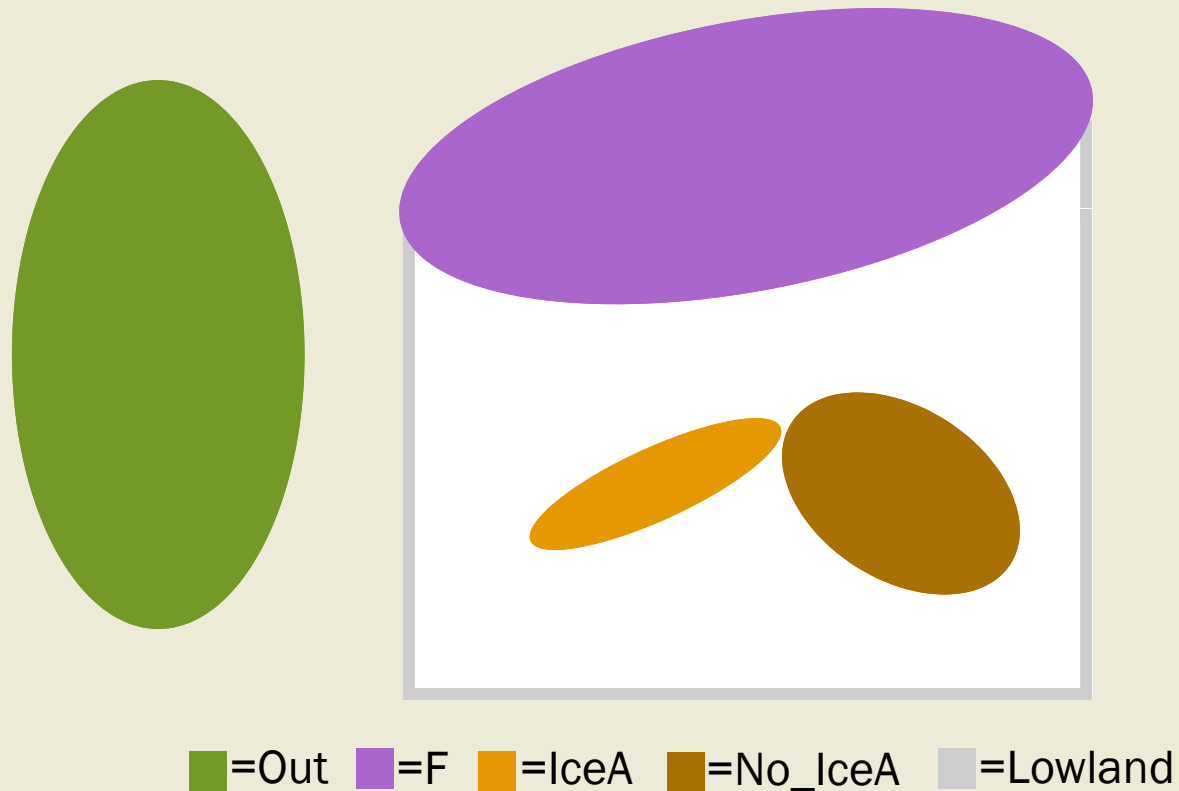
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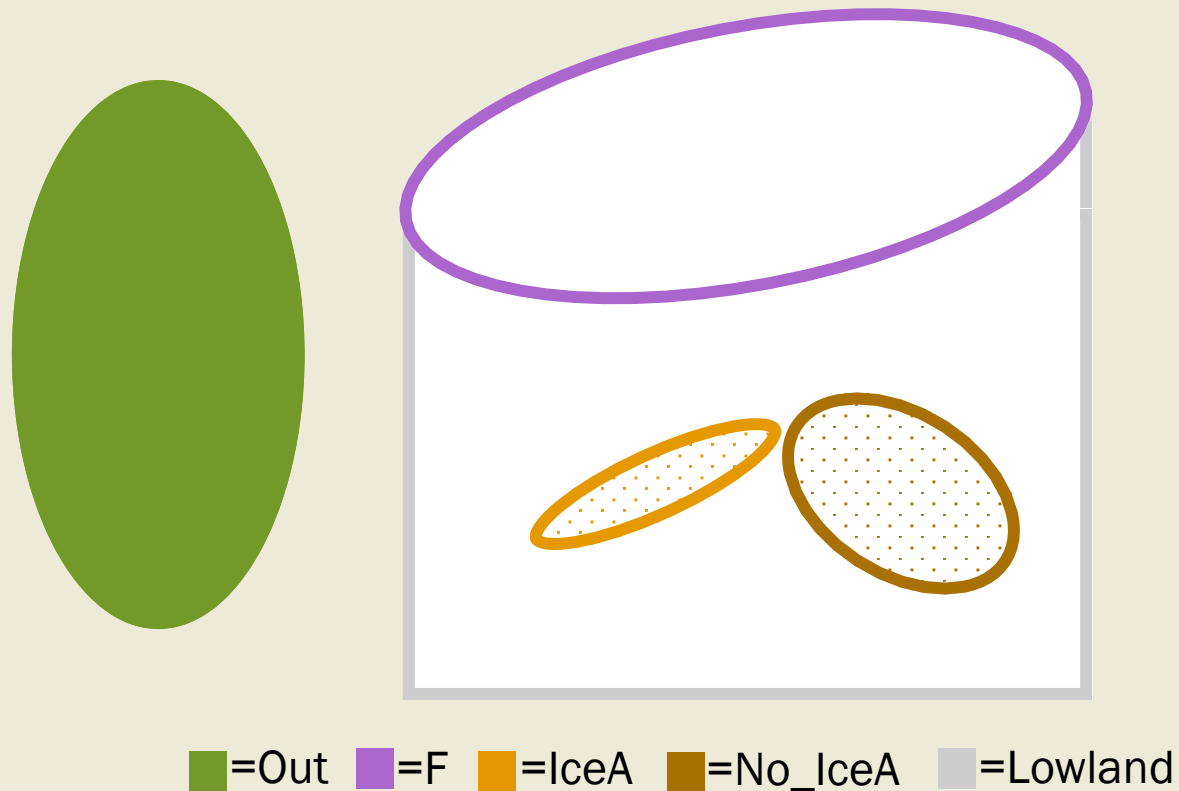
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Demographic scenarios

Alpine *Nunatak* scenario

LGM



LGM

- Lowland area not suitable
 - Too dry
- *Micro-refugia* in southern Alpine regions only

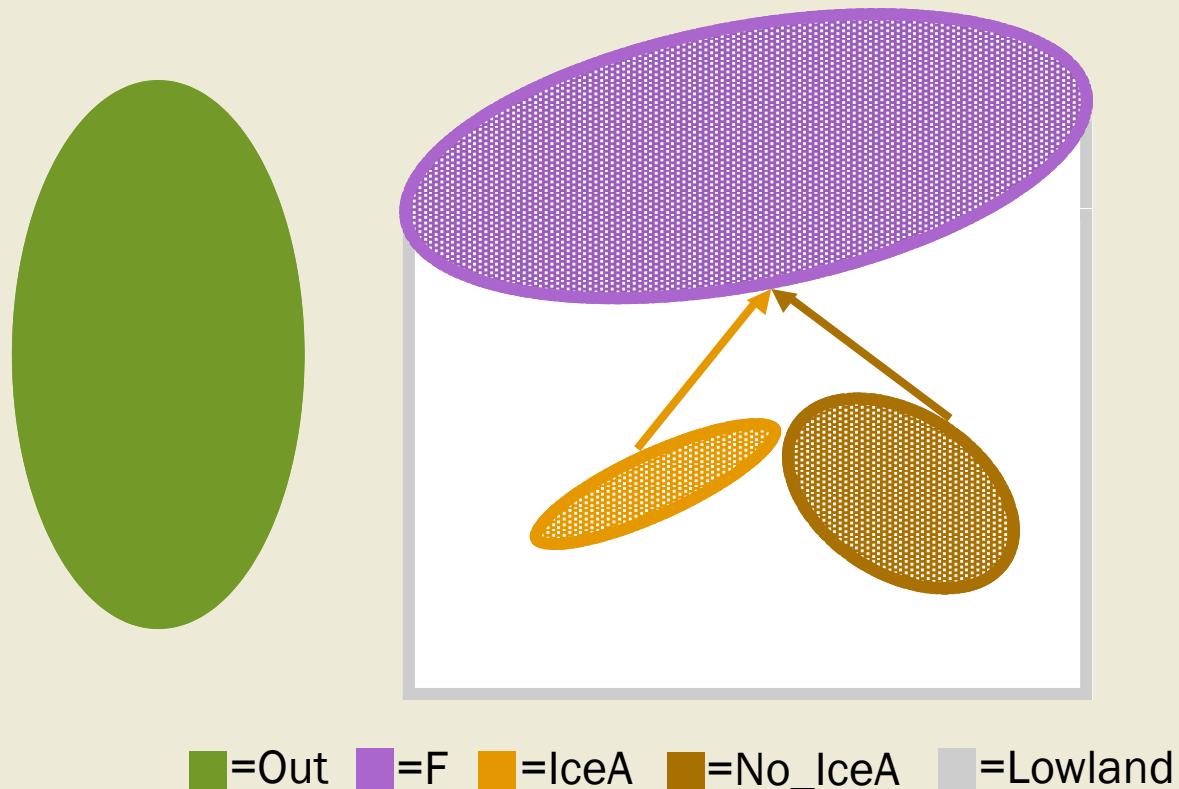
Onset

- Recolonization of Fennoscandia from those *refugia*
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Demographic scenarios

Alpine *Nunatak* scenario

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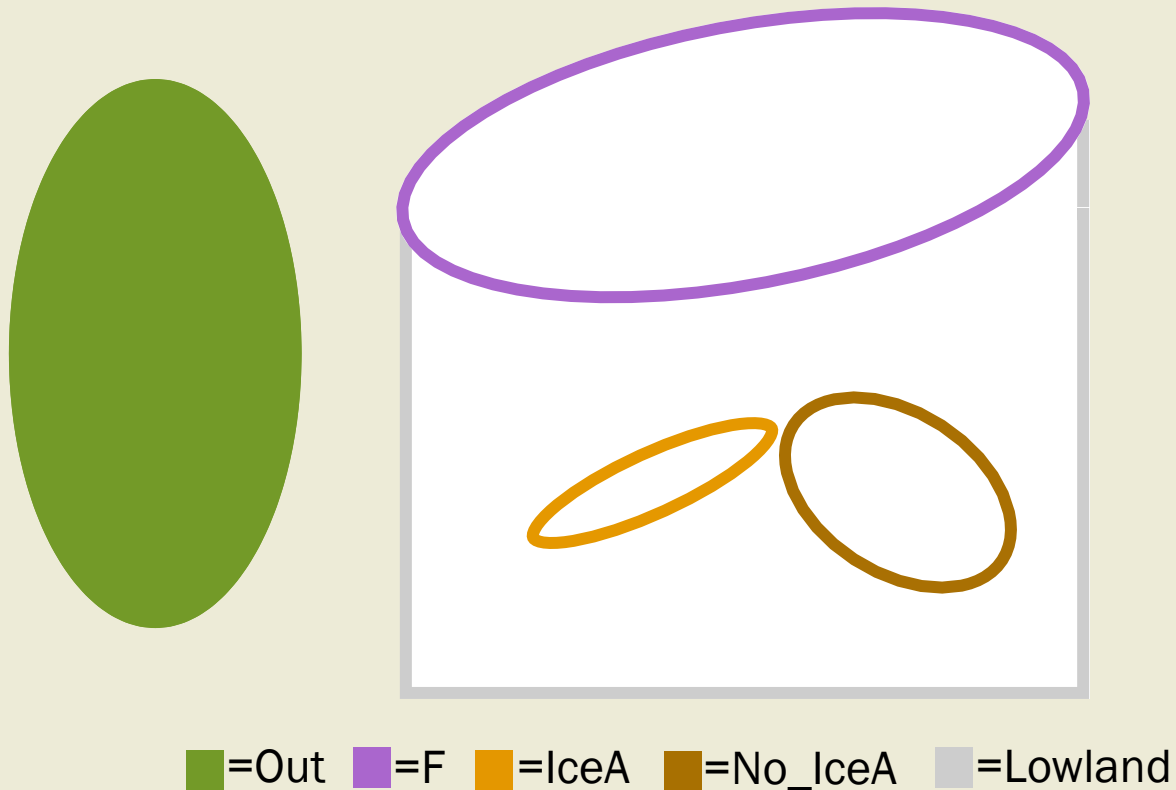
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Demographic scenarios

Out-of-Europe scenario

LGM



LGM

- No survival within the ice sheet
- Lowland area not suitable either
 - Too dry

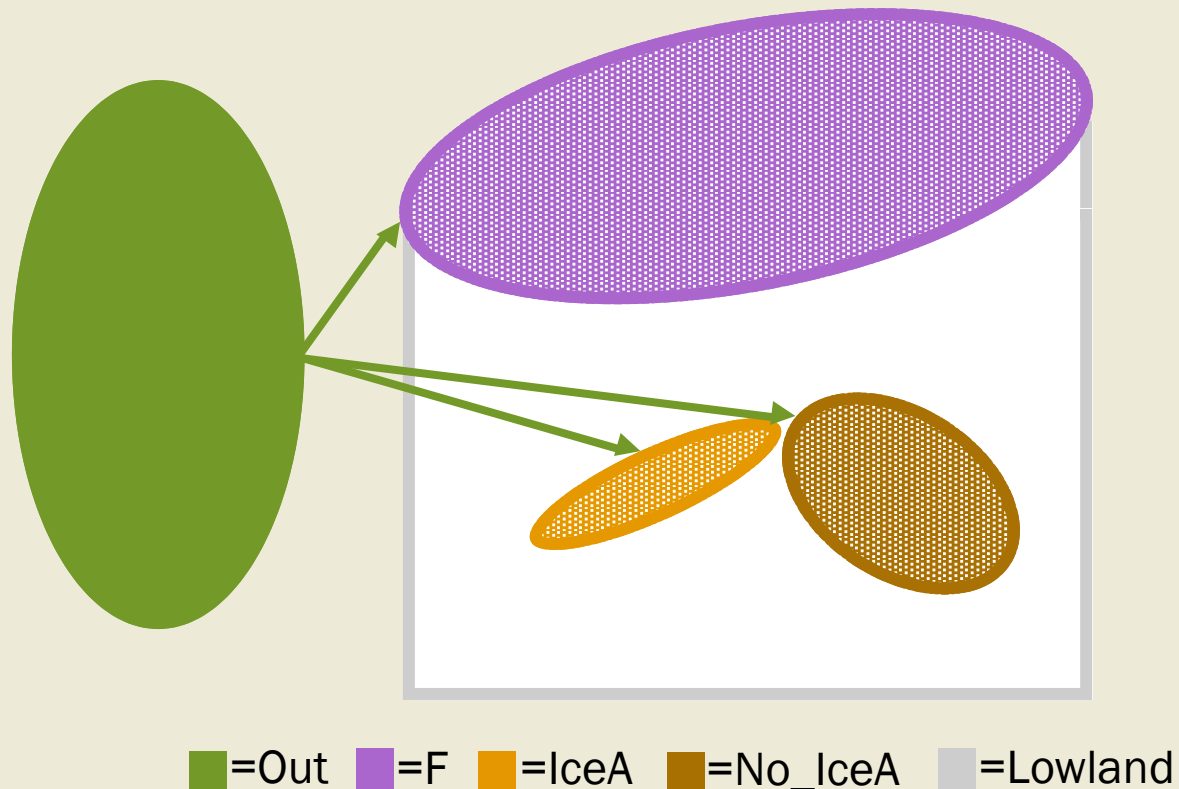
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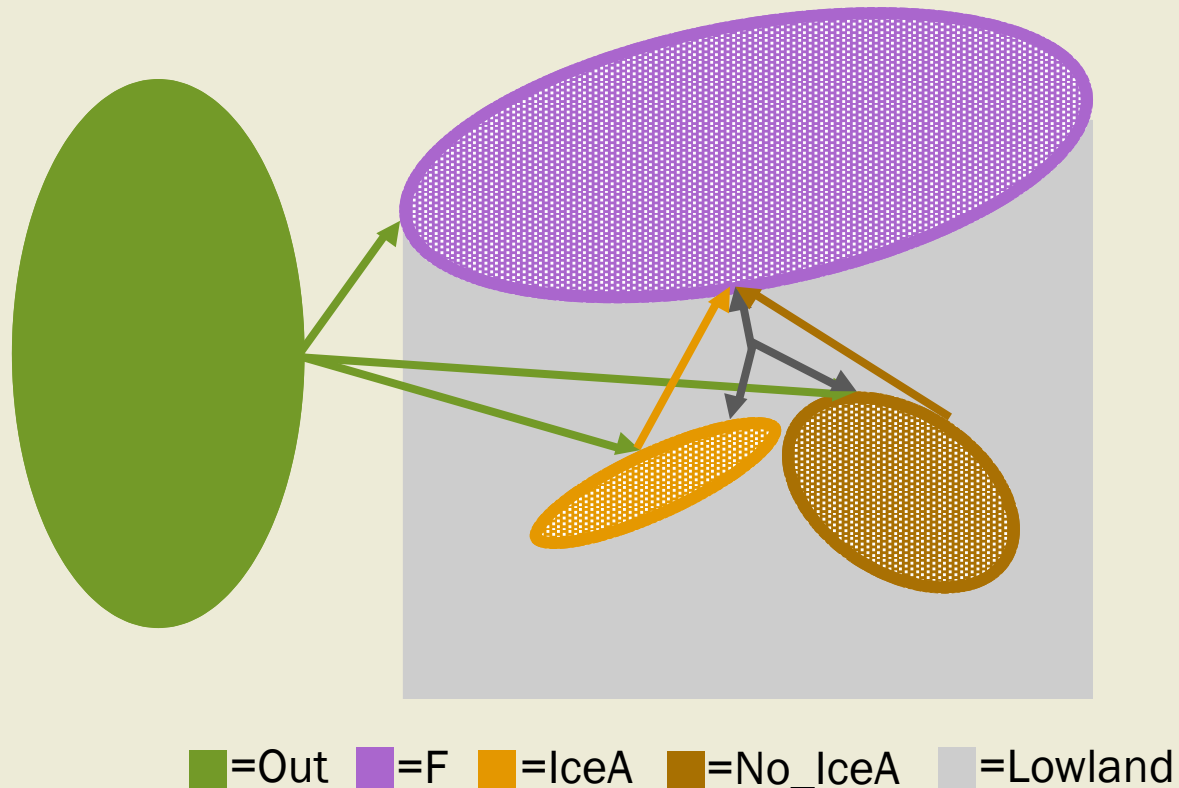
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Demographic scenarios

Composite scenario a

Onset



LGM

- Lowland areas suitable
- *In-situ* survival in micro-refugia within the ice sheet

Onset

- Migration rates from both Lowland and Out-of-Europe areas to Arctic-Alpine regions
- Migration rates from Alpine regions to Fennoscandia

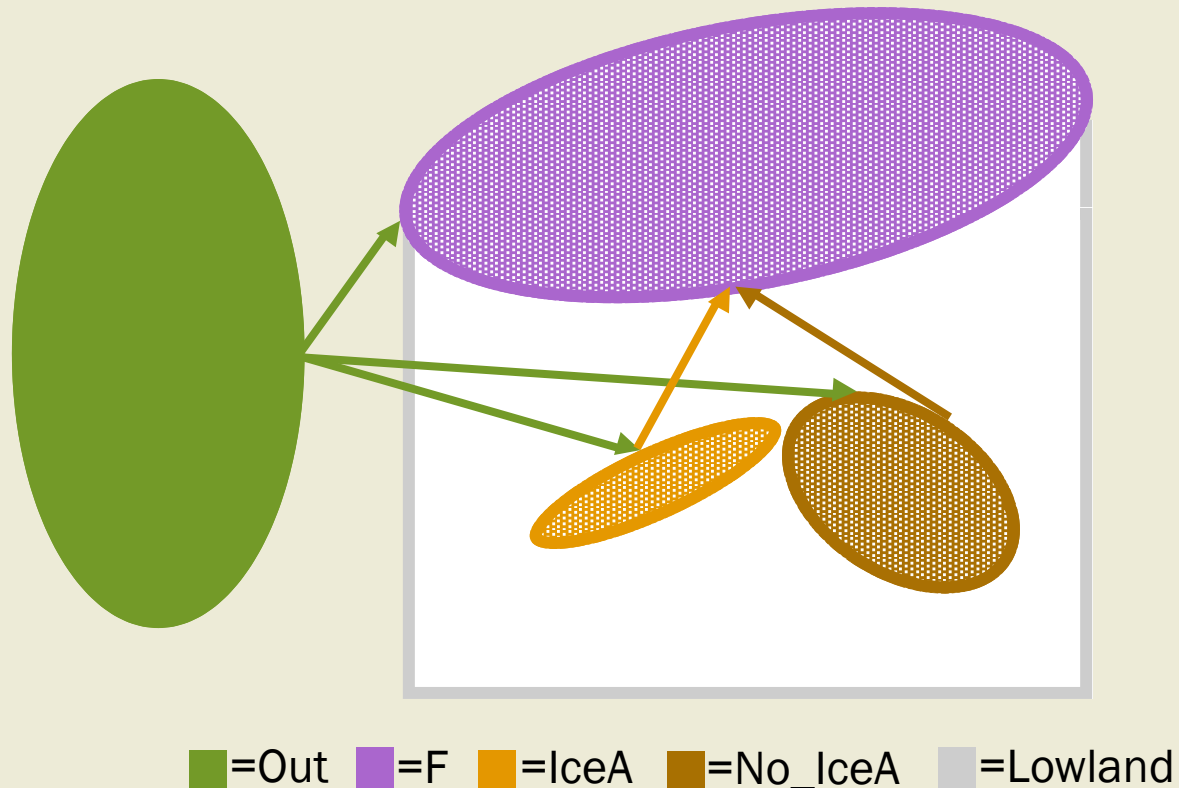
Present

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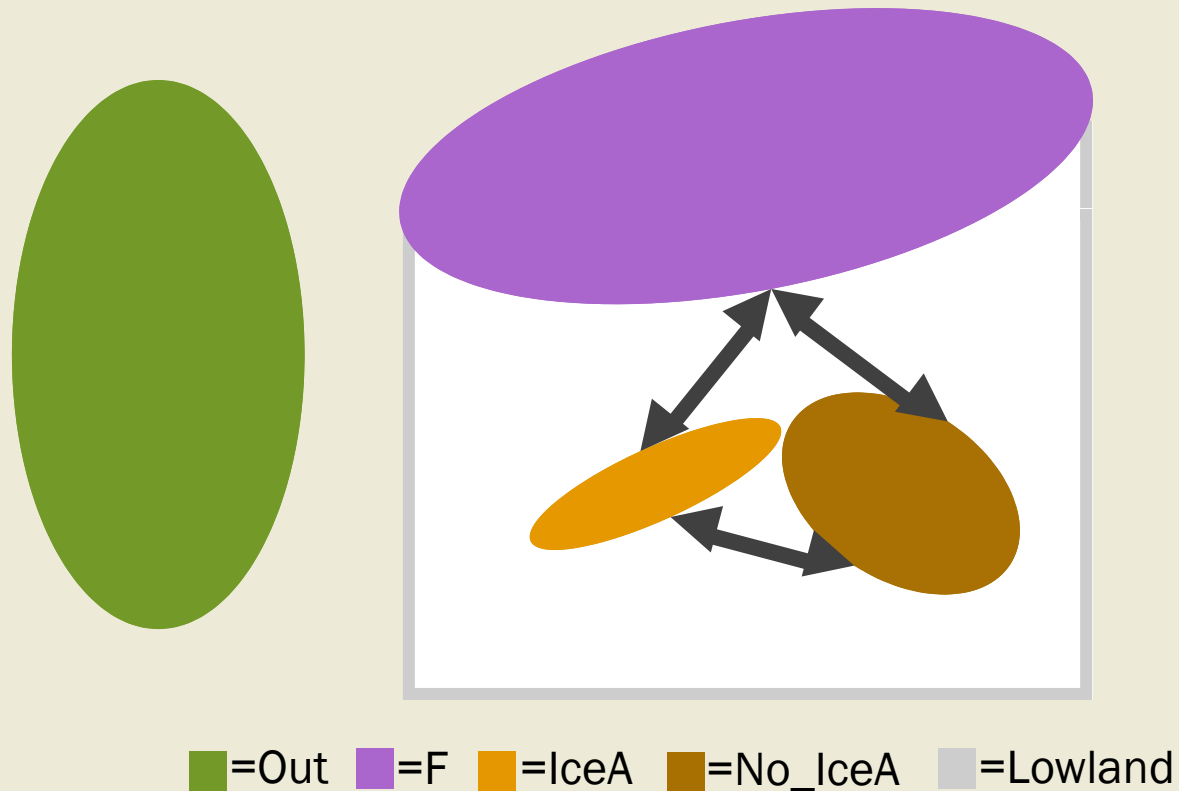
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- Migration rates from Out-of-Europe areas to Arctic-Alpine regions
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Demographic scenarios

Nul hypothesis (H0): Test for phylogeographic signal

Present



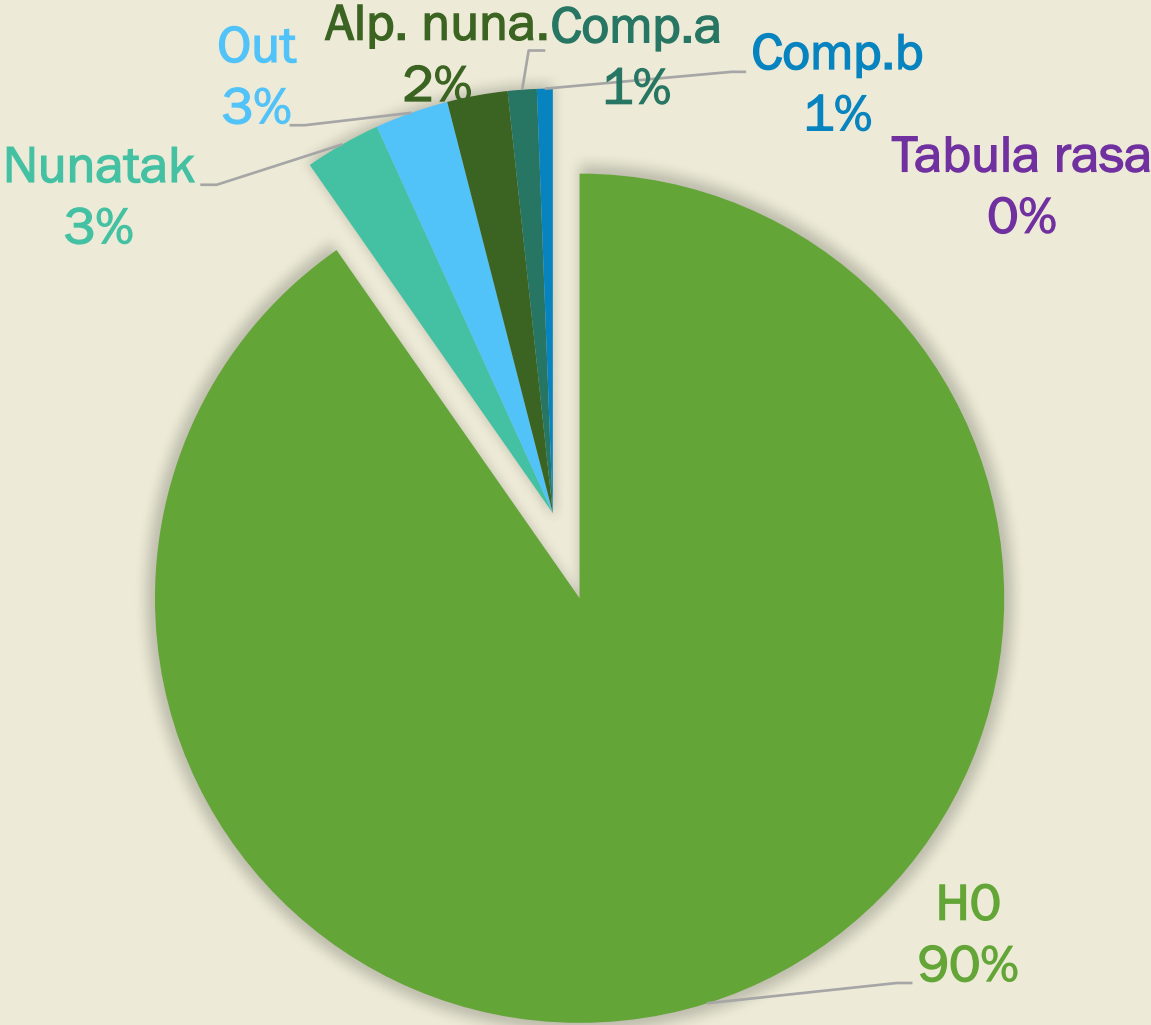
Present

- Whatever happened before, post-glacial migration rates within Europe erase any historical signal

Results and discussion

Posterior probability of each scenario

- *Timmia bavarica*



Best-Fit scenario : H0

⇒ Null hypothesis!

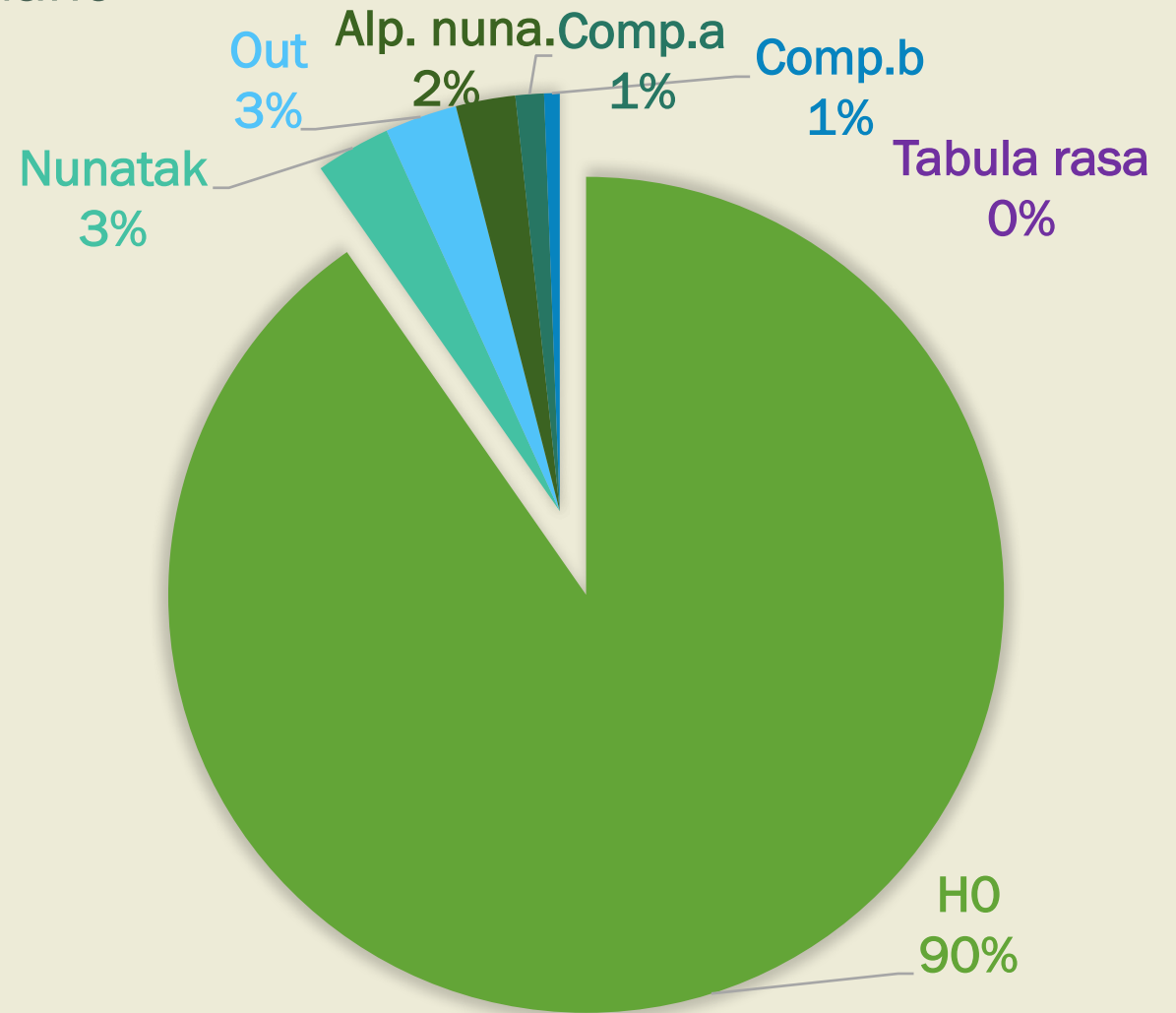
⇒ No phylogeographic signal in the data!

Tabula rasa : 0%

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Results and discussion

Posterior probability of each scenario

- *Amphidium lapponicum*

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H0
100%

Comp.b
0%

Out
0%

Nunatak
0%

Comp.a
0%

Alp. nuna.
0%

Tabula rasa
0%

Results and discussion

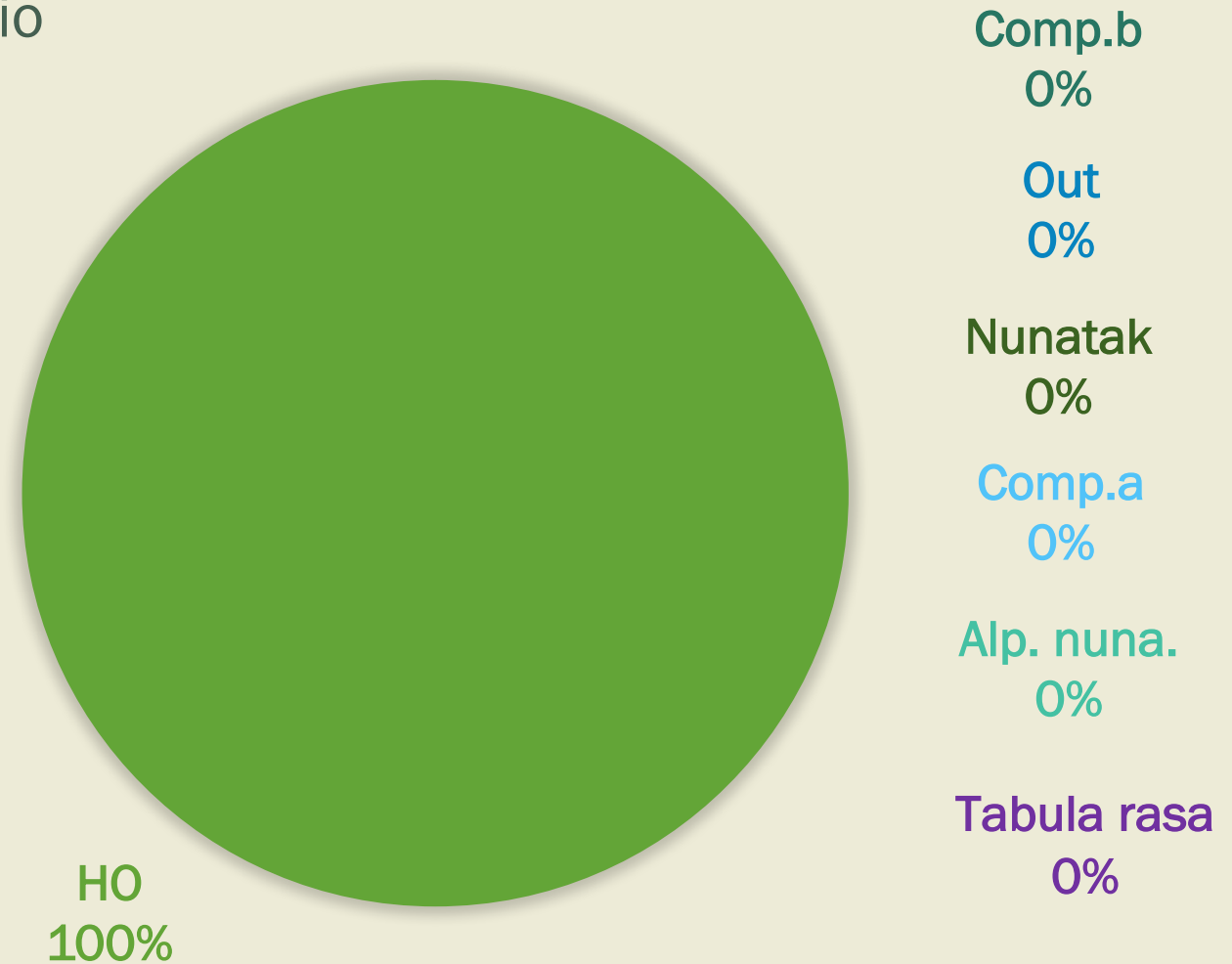
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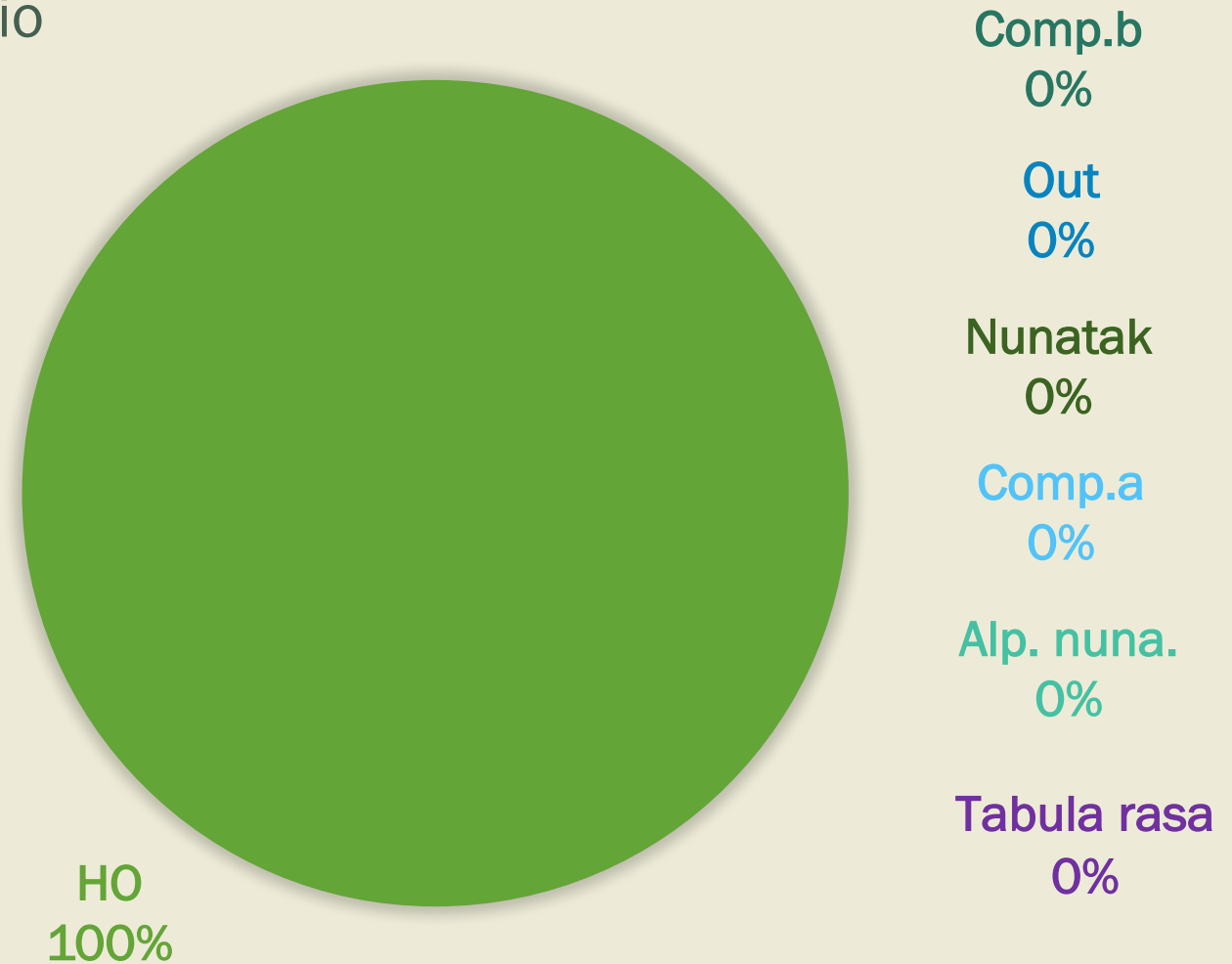
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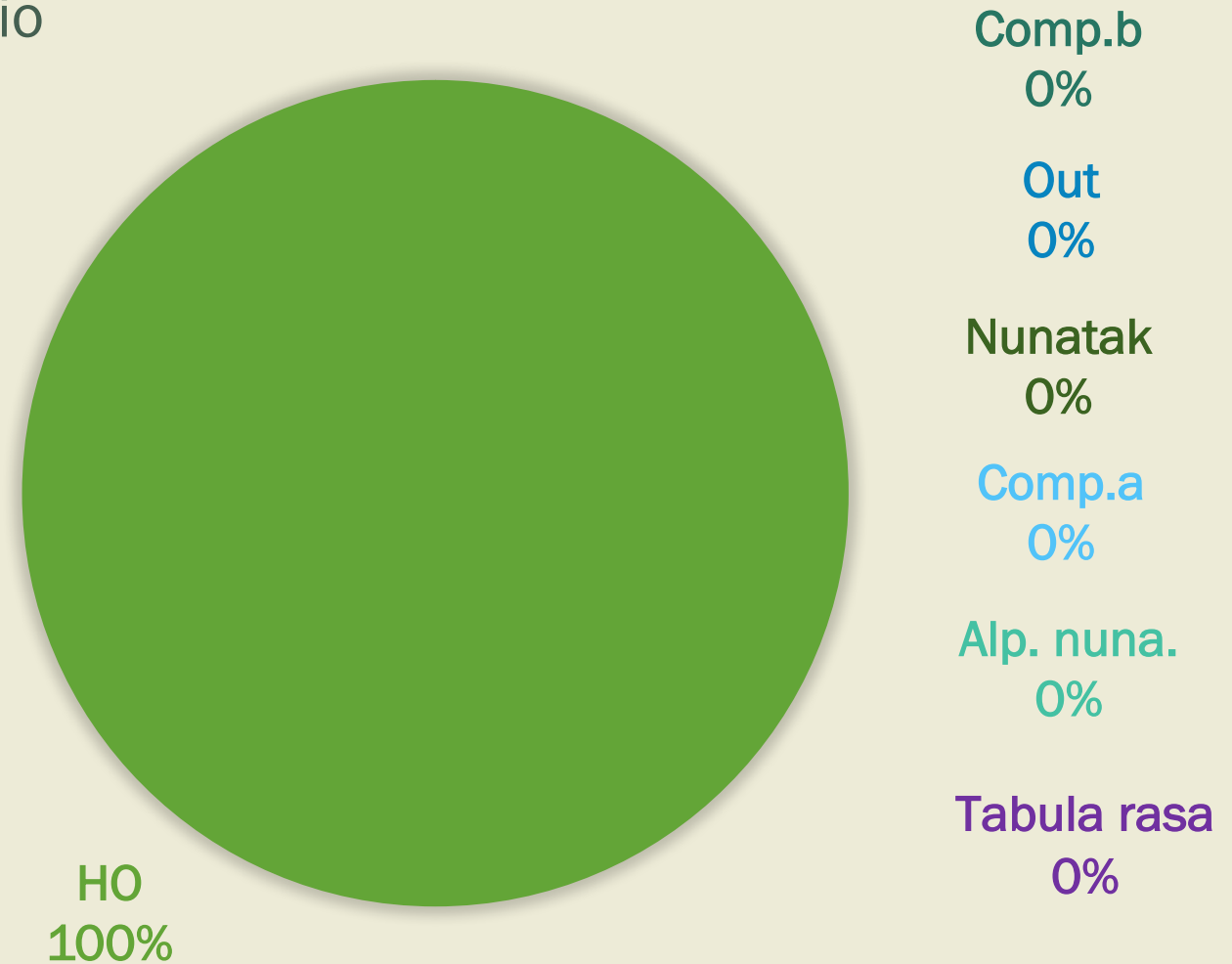
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H0 is the best-fit scenario!

- Unexpected result
- Actual migration rates within Europe erase any trace of historical signal

⇒ Highlights the high dispersal capacities of bryophytes

Consequence

- Impossible to retrace the biogeographic history of the Arctic-Alpine mosses in Europe...

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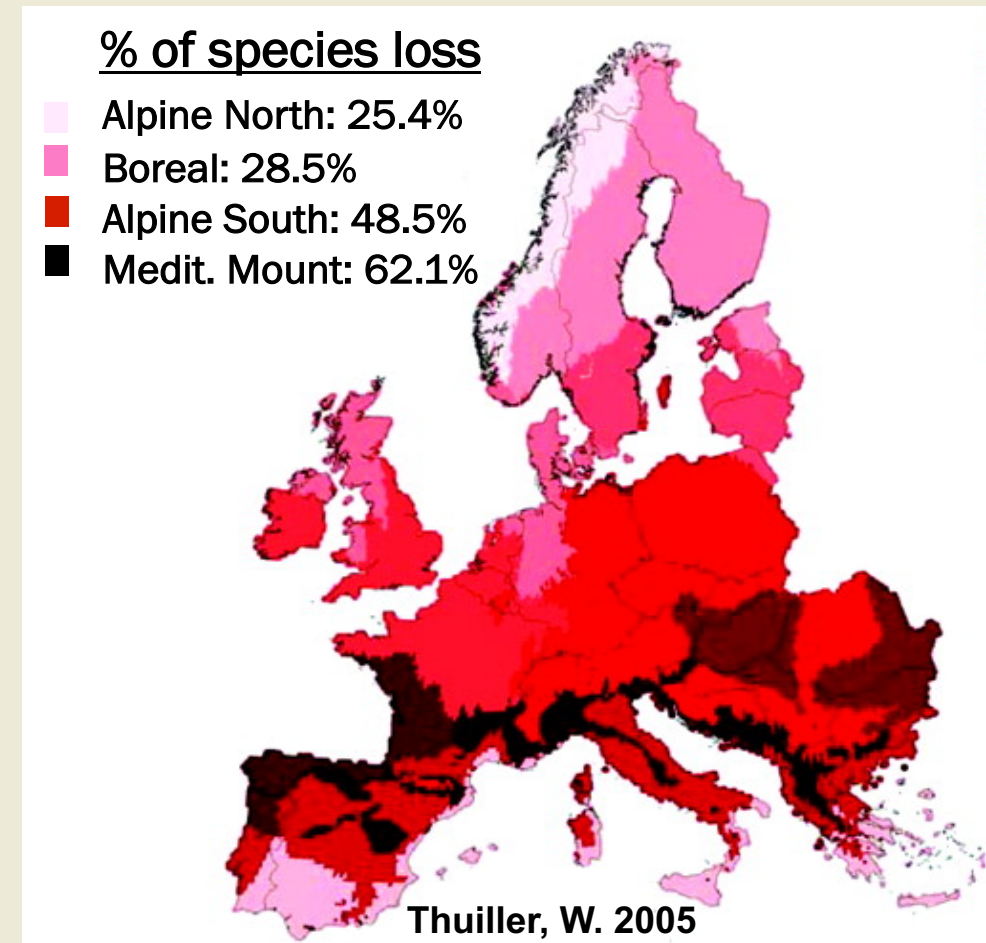
Conclusion and perspectives

Arctic-Alpine populations highly endangered

- In the context of climate change
- Especially Alpine populations
 - Small already
 - By 2080, 48.5% of the Alpine plant species will be lost against 28.5% for the Arctic ones

BUT : Great news!

- Alpine populations should easily find refuge in Arctic populations
 - Thanks to migrations and high dispersal capacities of Bryophytes!



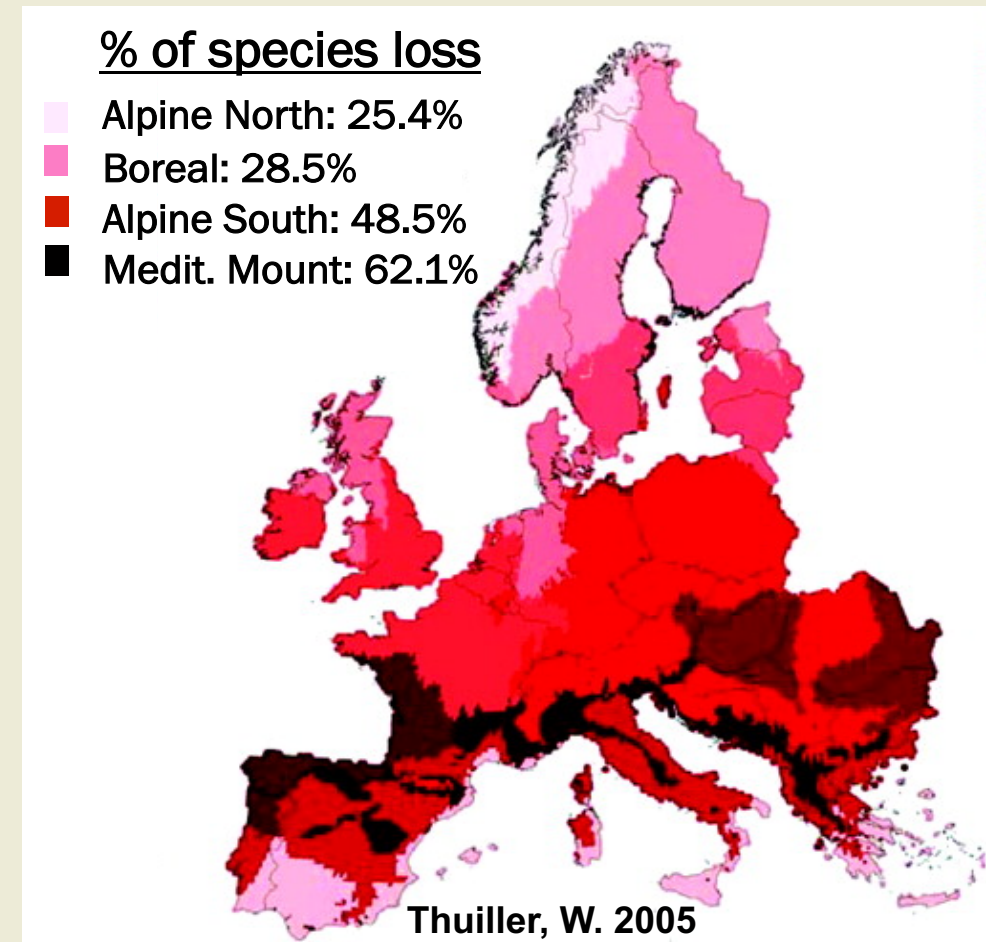
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THANK YOU FOR YOUR ATTENTION!

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