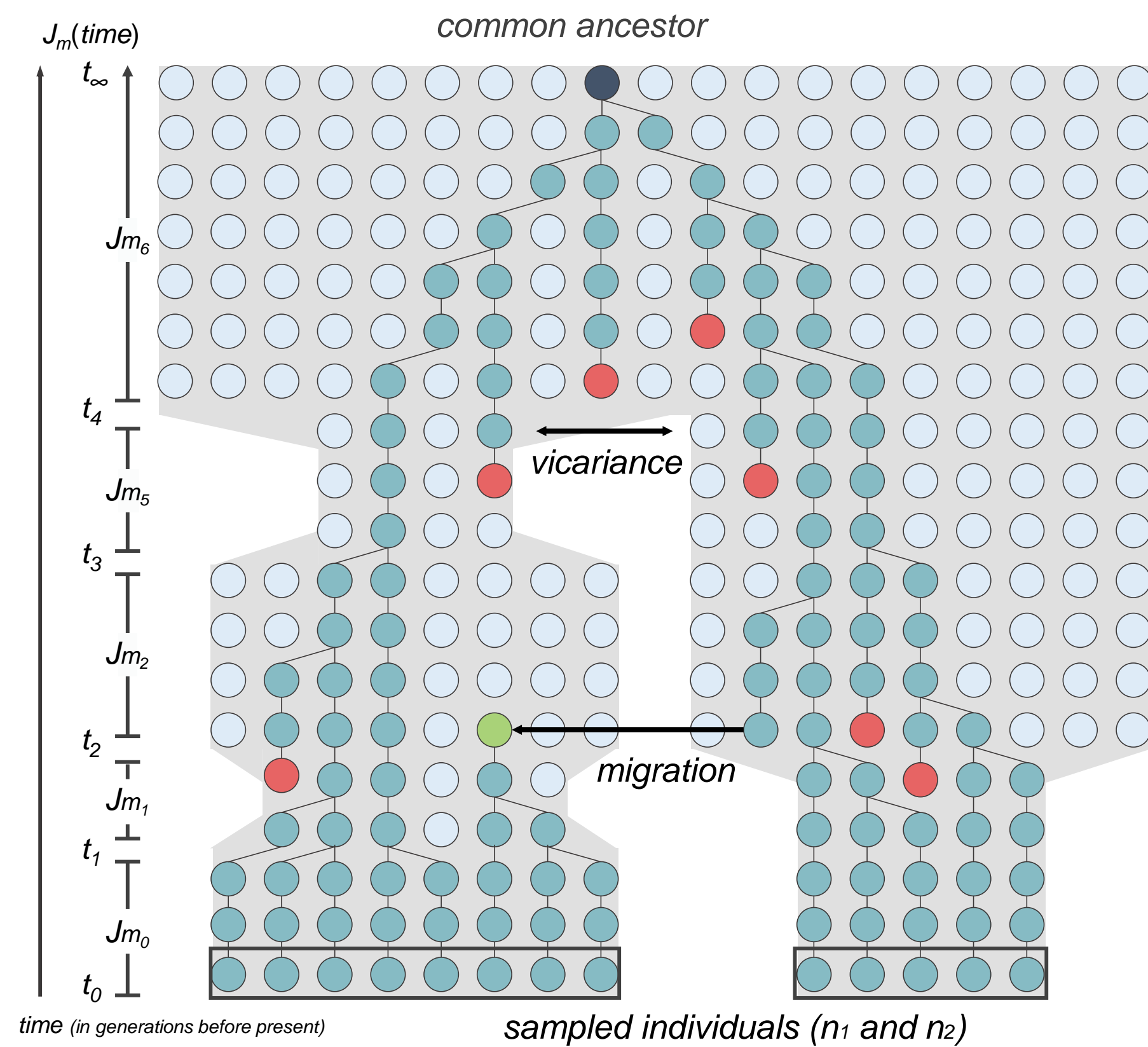


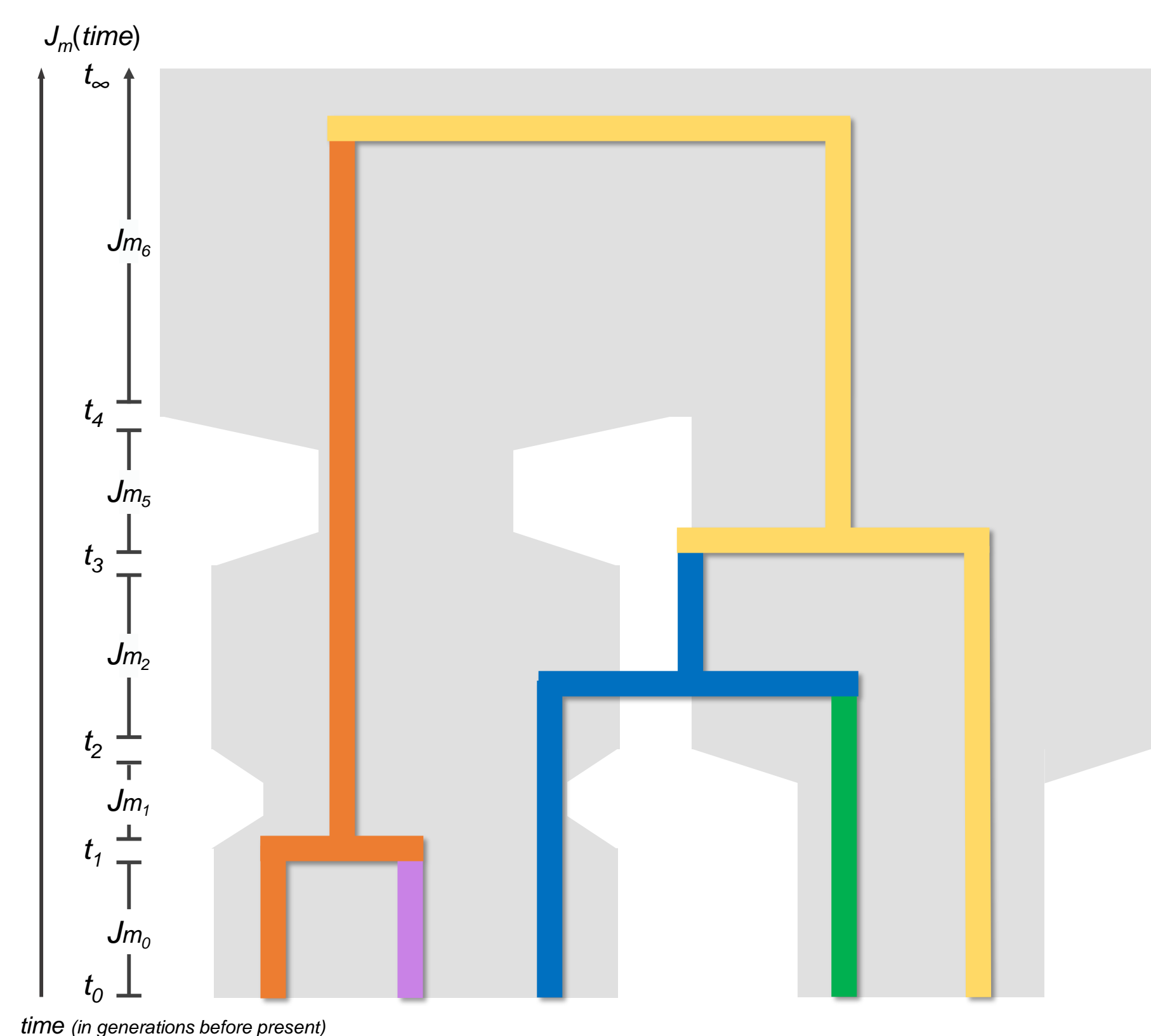
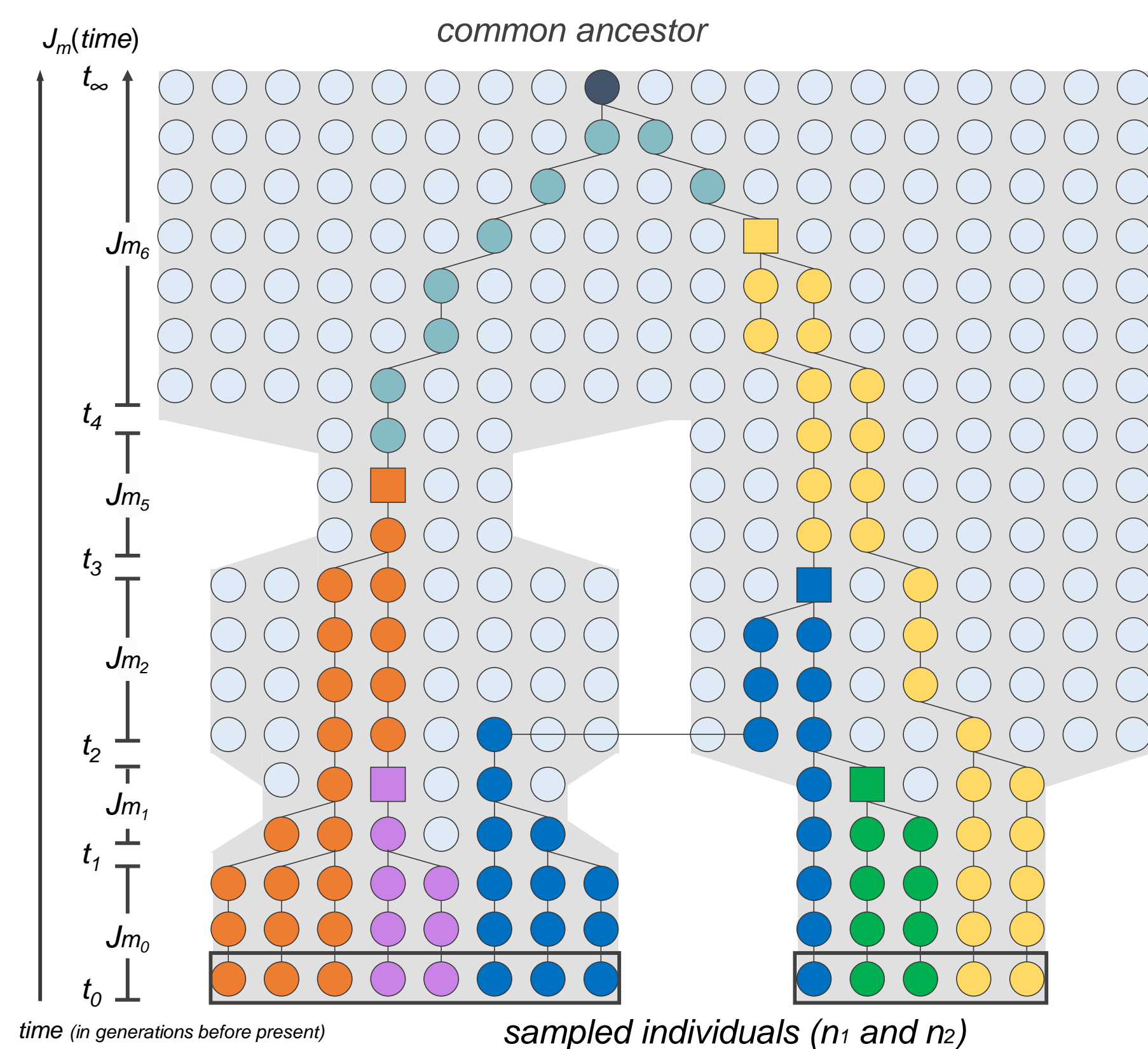
## Biogeographic history of individuals



## Model parameters

- The size of the different assemblages in the past (*deme\_sizes*) along with the corresponding dates (*changetimes*)
- The migration history linking demes (*migr* and *migr\_times*)
- The vicariance history linking demes (*splits*)
- The point mutation rate (*mu*)
- The speciation model (*smodel*, *tau*)

## Evolutionary relationships of species



From these users can derive a wide range of summary statistics of interest (species richness, Shannon's or Simpson's diversity indexes, phylogenetic diversity metric etc.)

3 2 4 2 2  
species relative abundances

Observations