



Model misspecification and overestimation of phylogenetic root age in linguistic and biological datasets

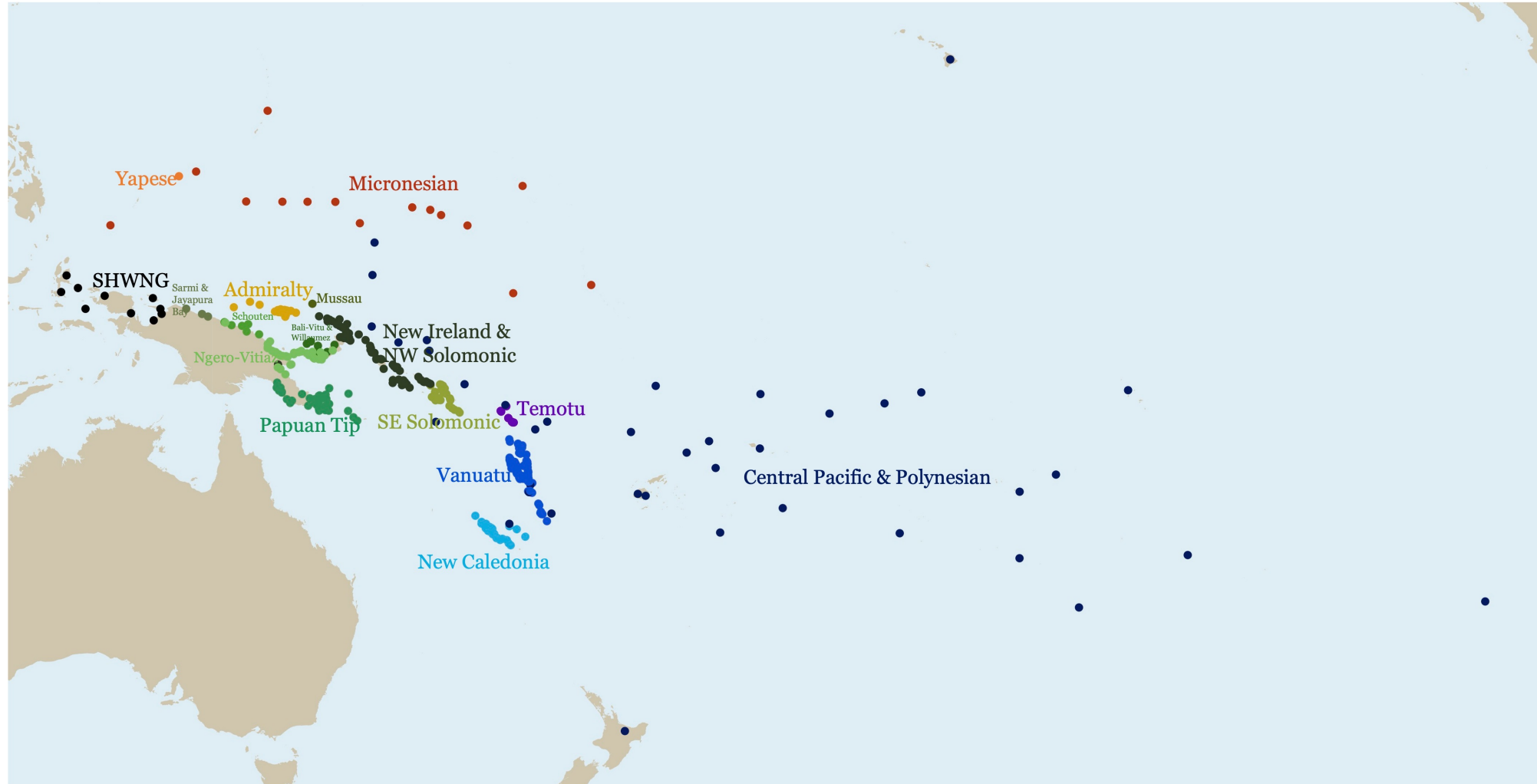
Benedict King, Aymeric Hermann, Mary Walworth, Simon Greenhill & Russell D Gray

Talk outline

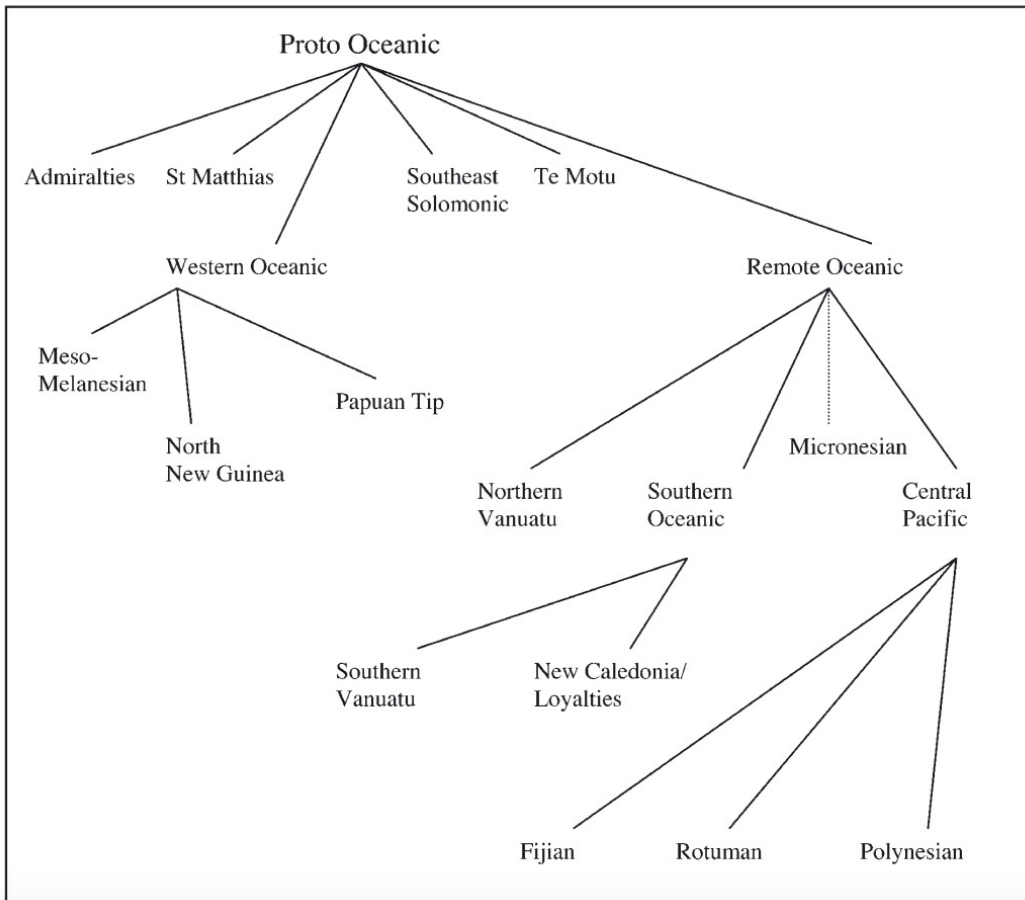


Phylo-purgatory

Oceanic languages



Aims

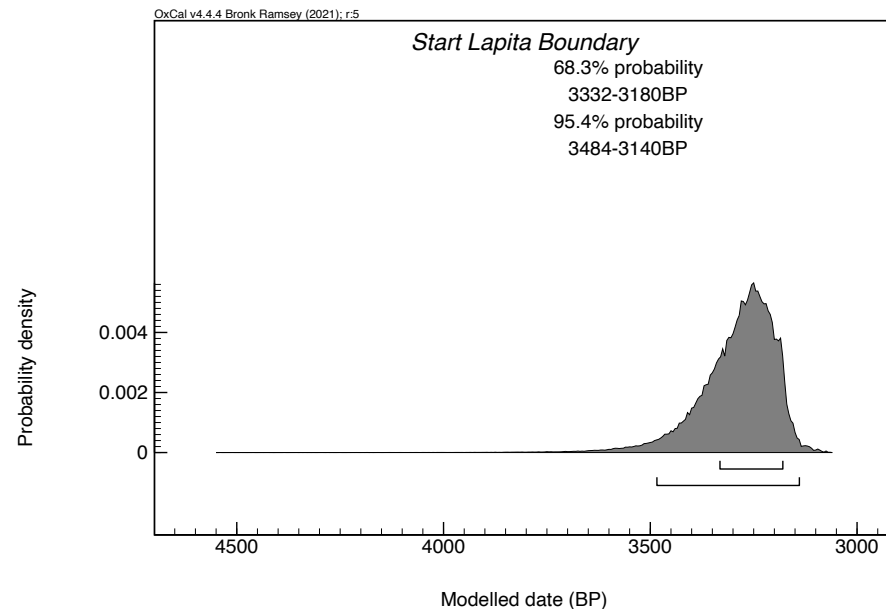


“ Despite the relatively clear evidence for Oc itself, which has been known at least since Dempwolff (1927), the higher-level subgroups of Oc have been extremely elusive ” Blust (1998)

- Sort out subgrouping
- Timing and order of migration

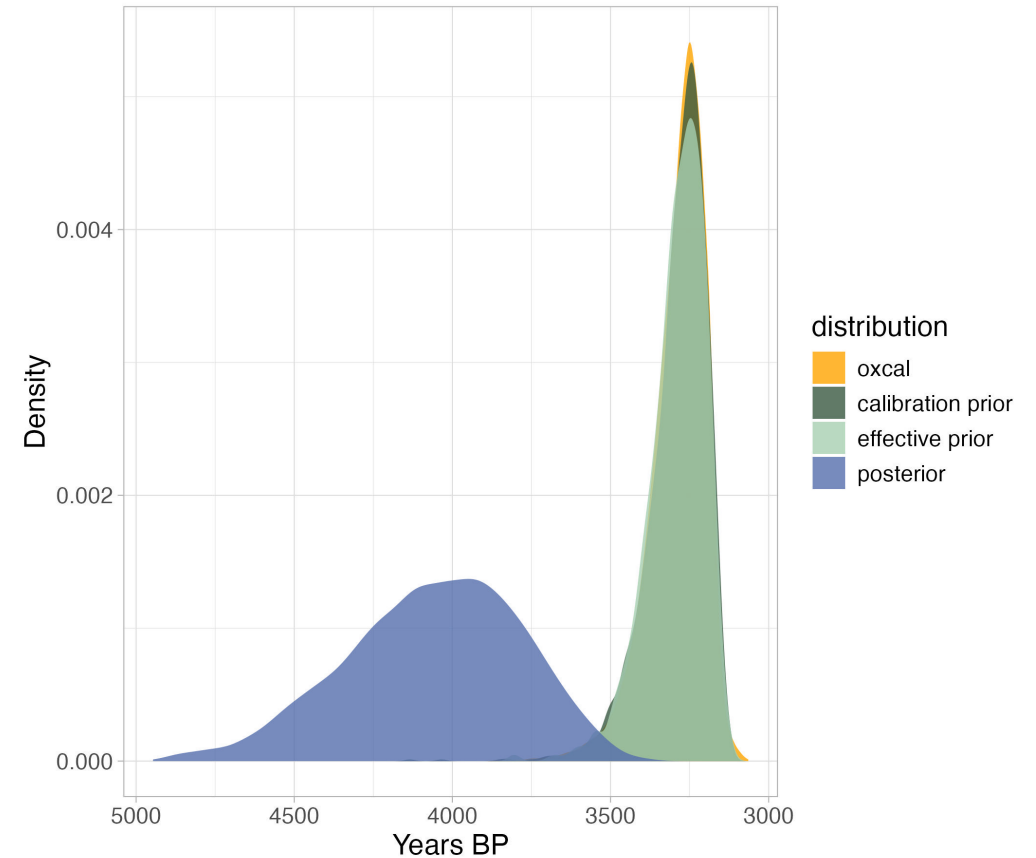
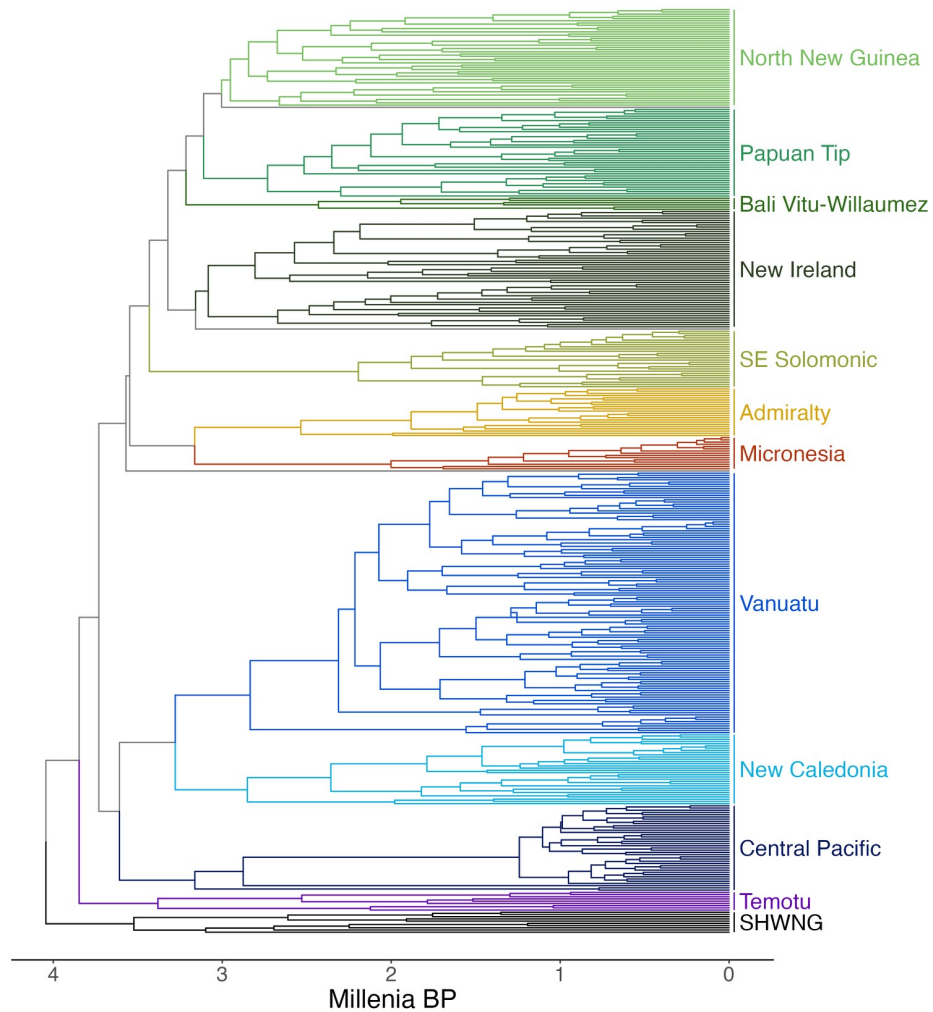
Oceanic phylogeny set-up

Austronesian Basic Vocabulary Database



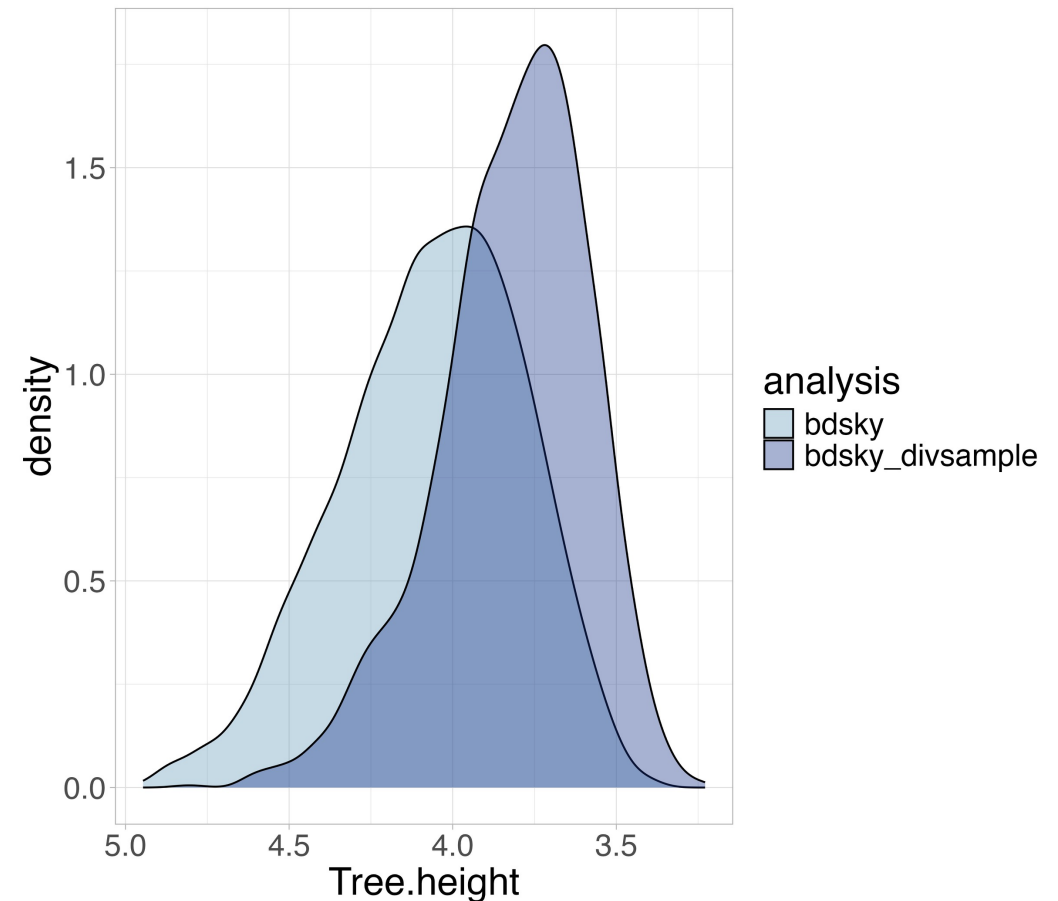
- Basic vocabulary data for 417 Oceanic languages
- 3 node calibrations based on archaeological dates
- Analysed in beast2 using the Birth-Death Skyline model

Oceanic phylogeny results



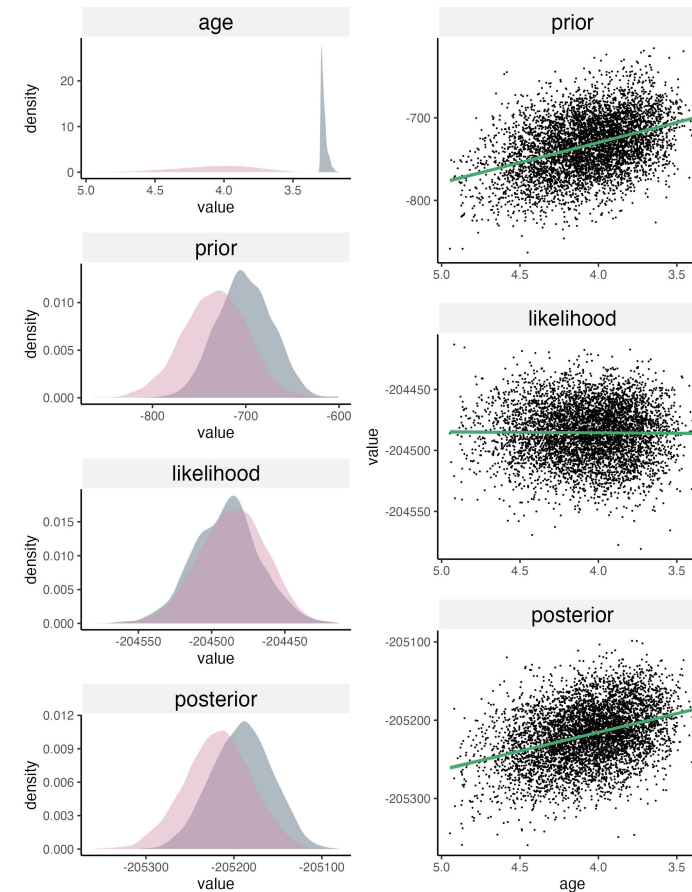
Deep Root Attraction

- Can be caused by diversified sampling
- Often the effective prior does not match the calibration density
- Does not explain Oceanic results

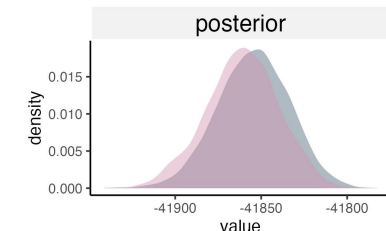
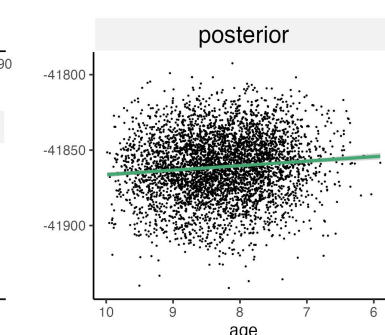
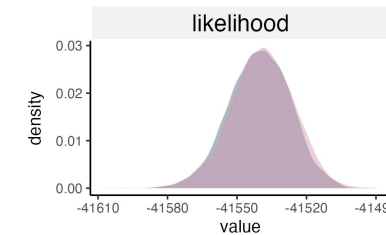
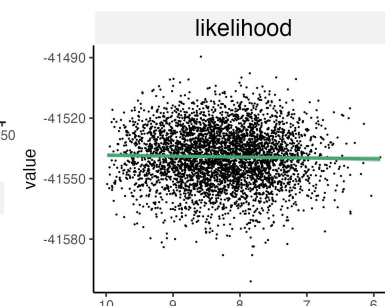
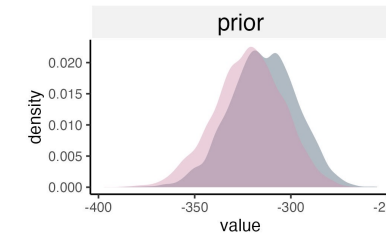
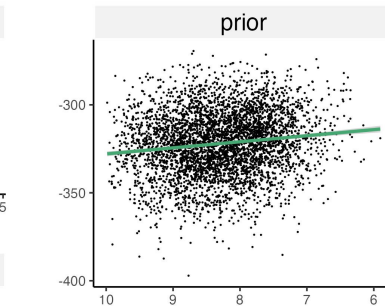
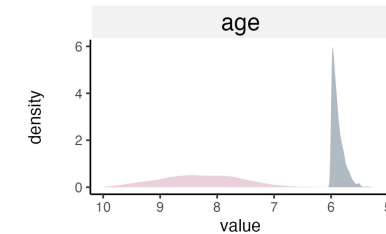
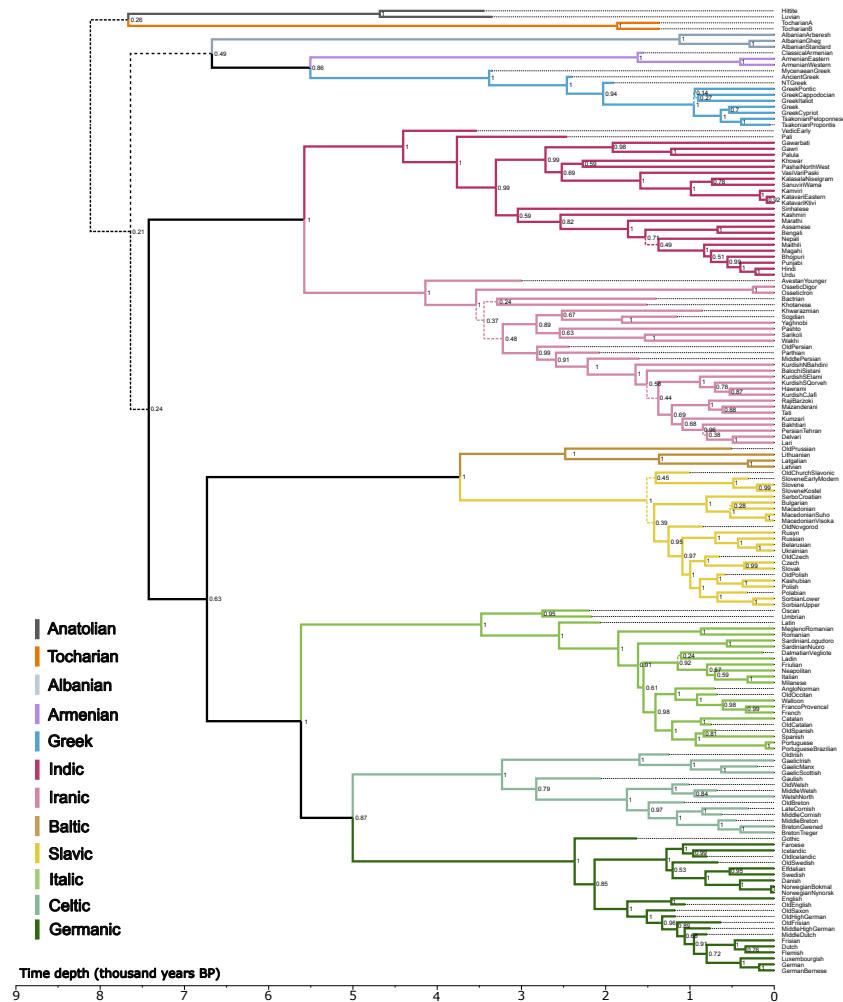


Does the data support older ages?

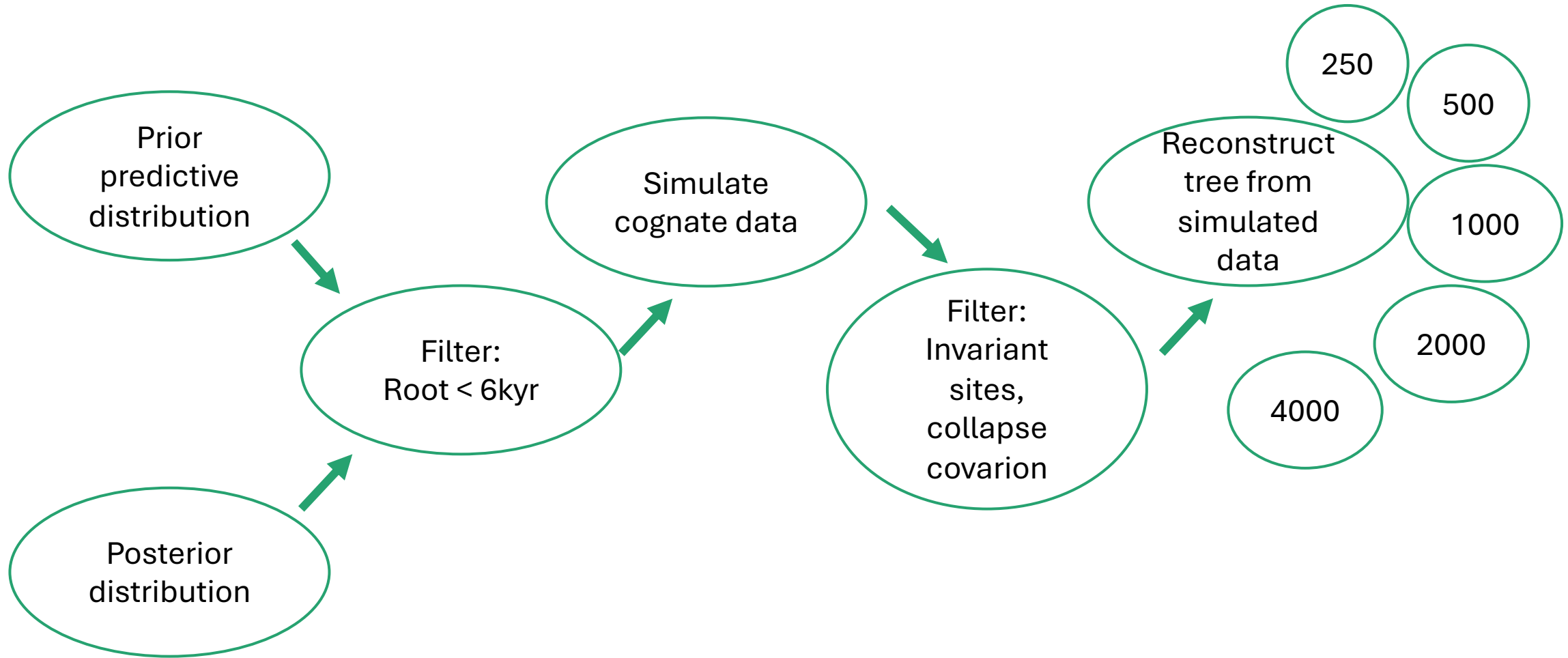
- Constrain a maximum age of 3300 BP
- Likelihood is identical regardless of age



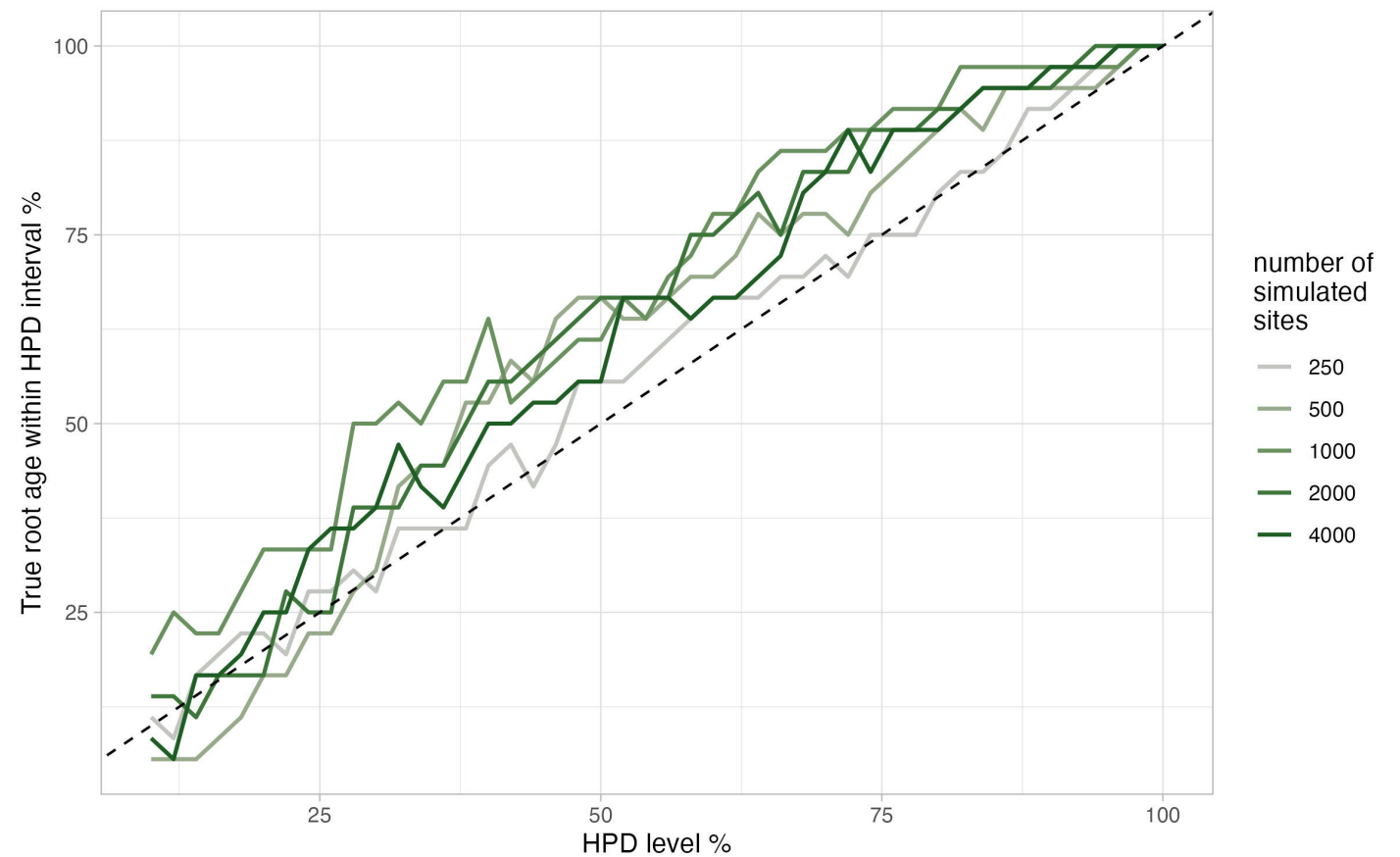
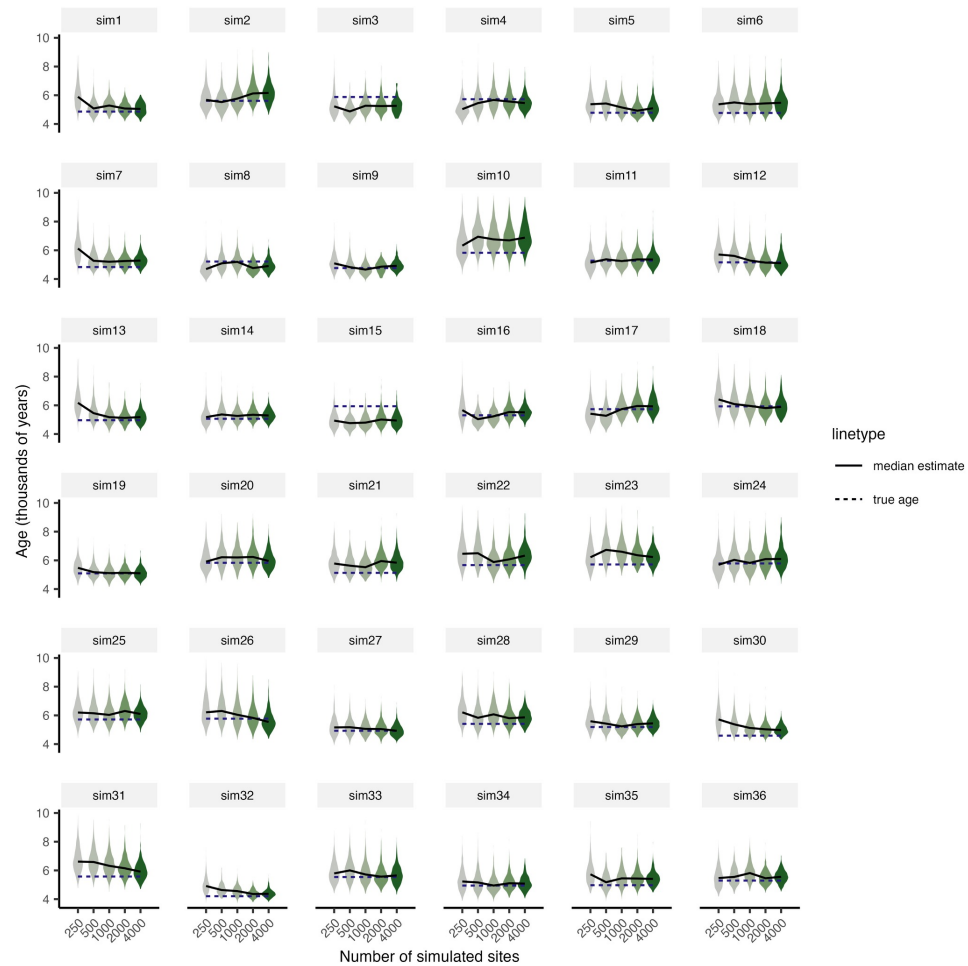
What about other language families?



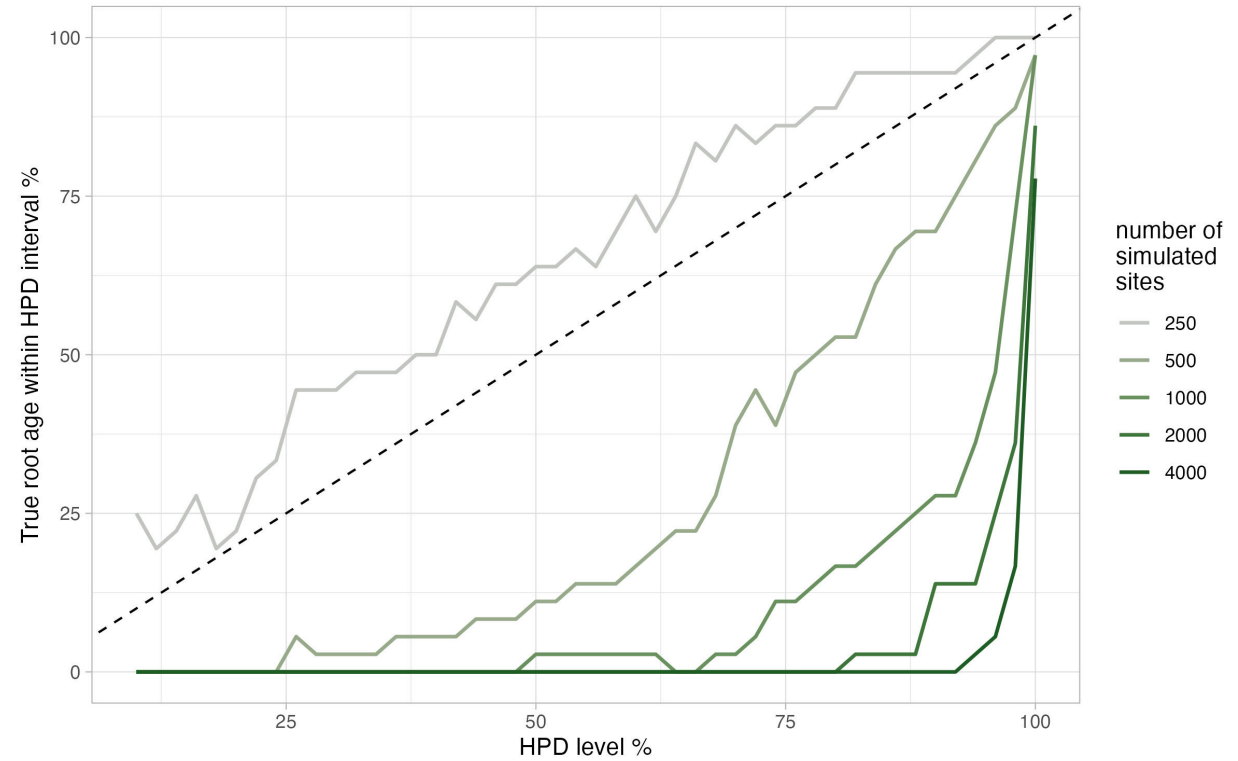
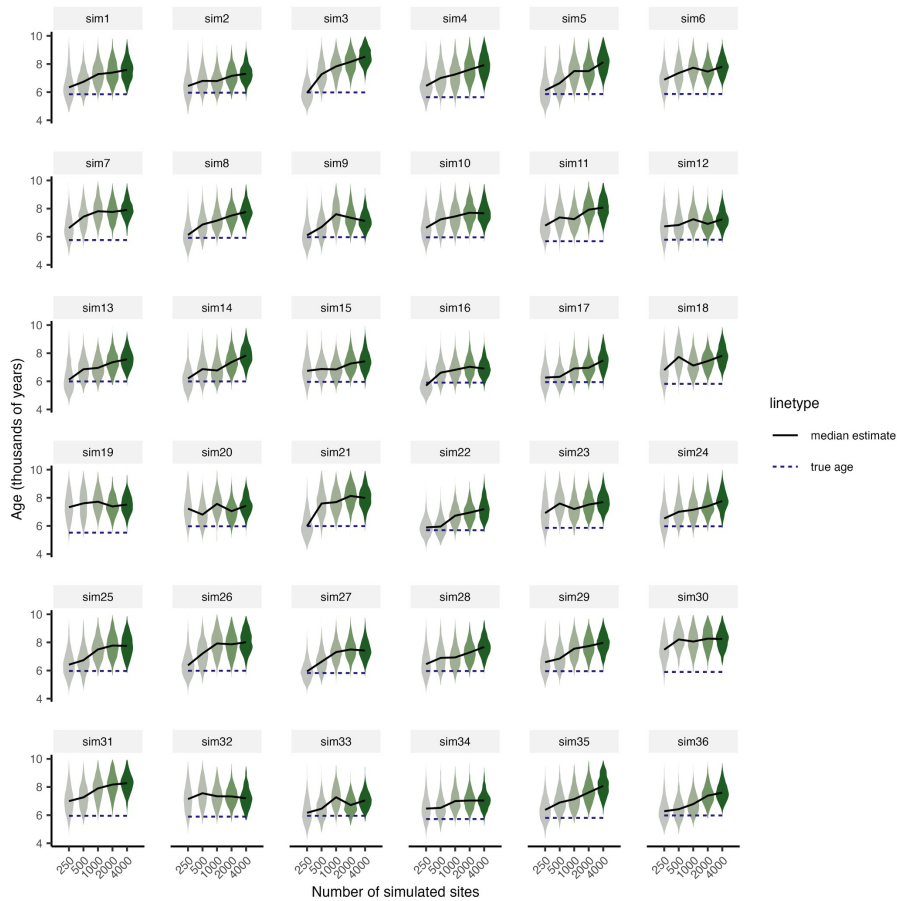
Can the methods get young trees?



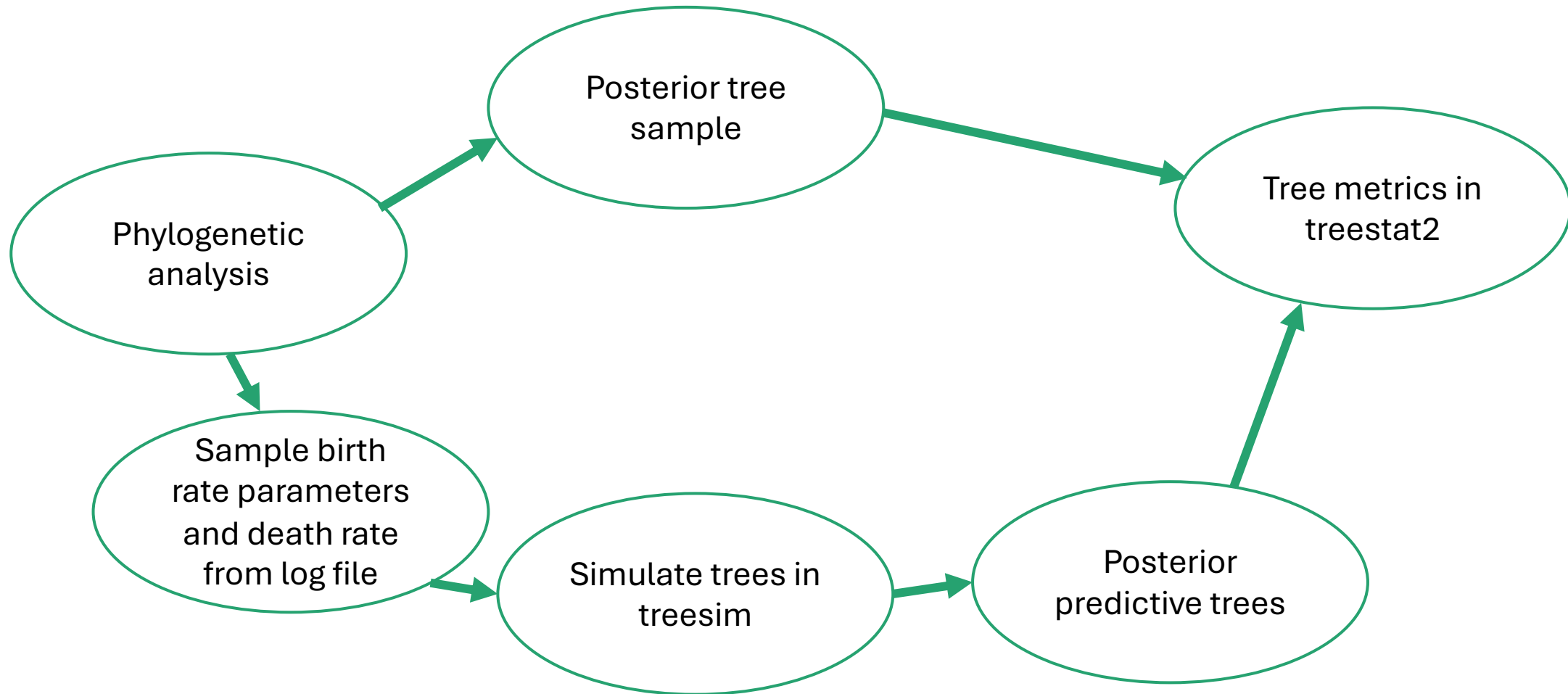
Results – trees from the prior



Results – trees from the posterior

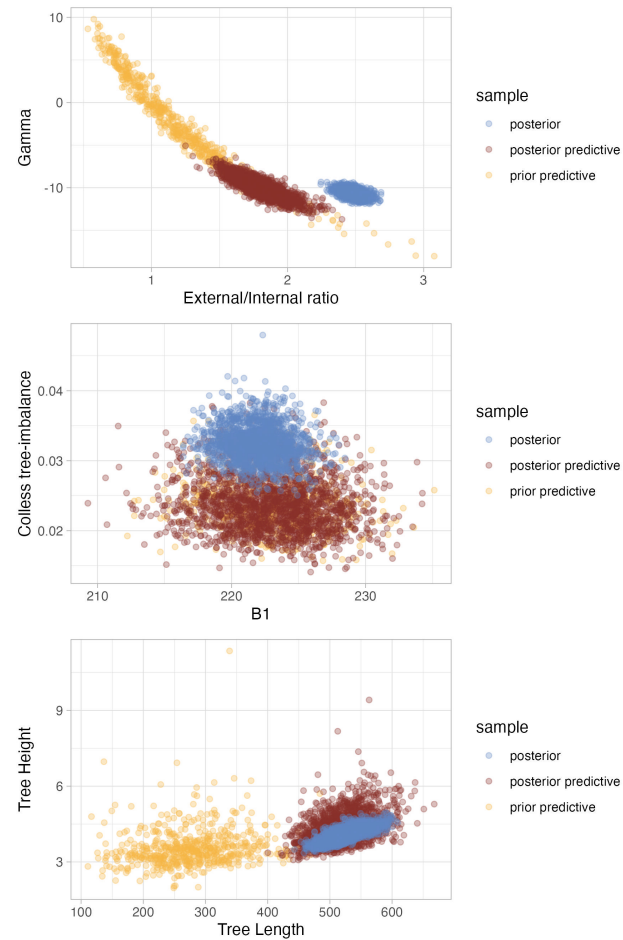


Posterior tree fit

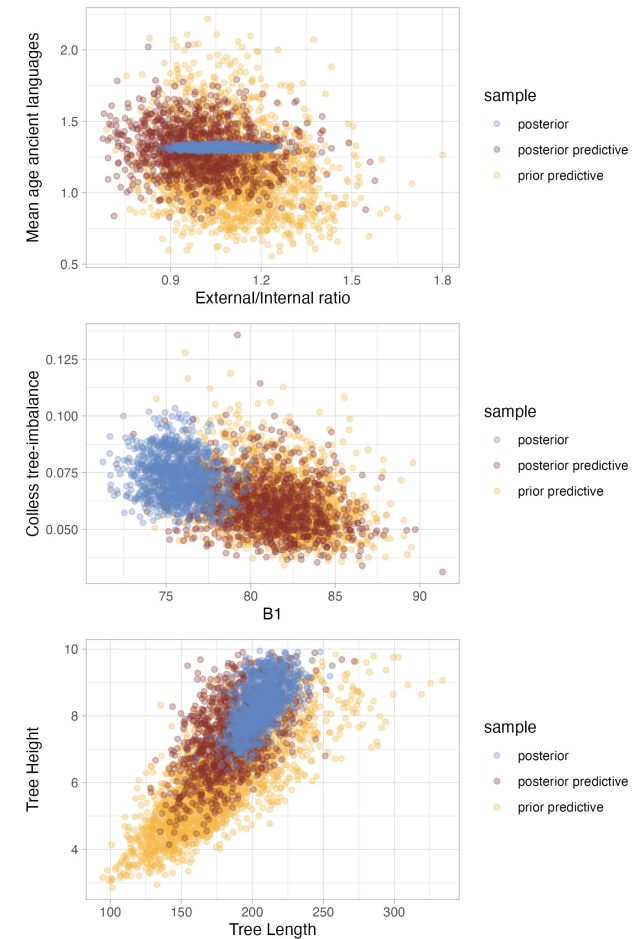


Posterior predictive results

Oceanic



IE



Is this a linguistics problem?

“*It’s the way the cognates are collected*”

“*It must be the covarion model*”

“*Biological models don’t work in Linguistics*”

Do biological models work in Biology?



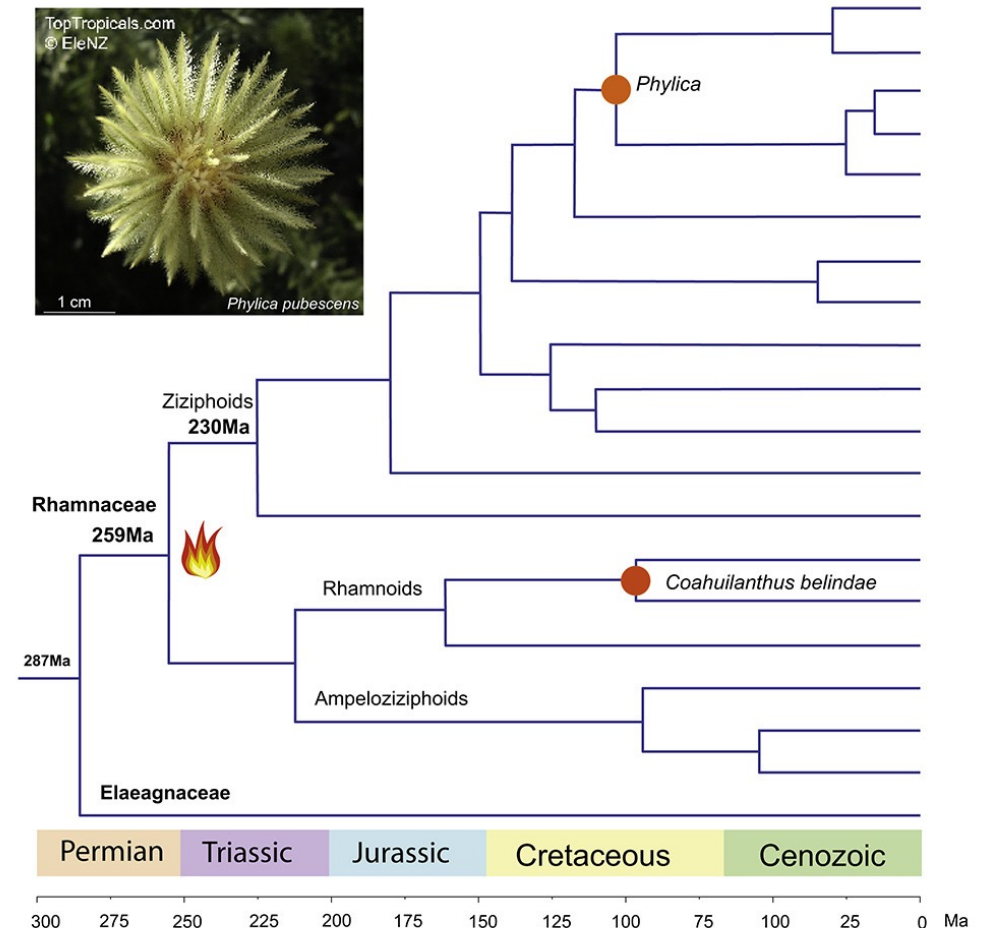
iScience

CellPress
OPEN ACCESS

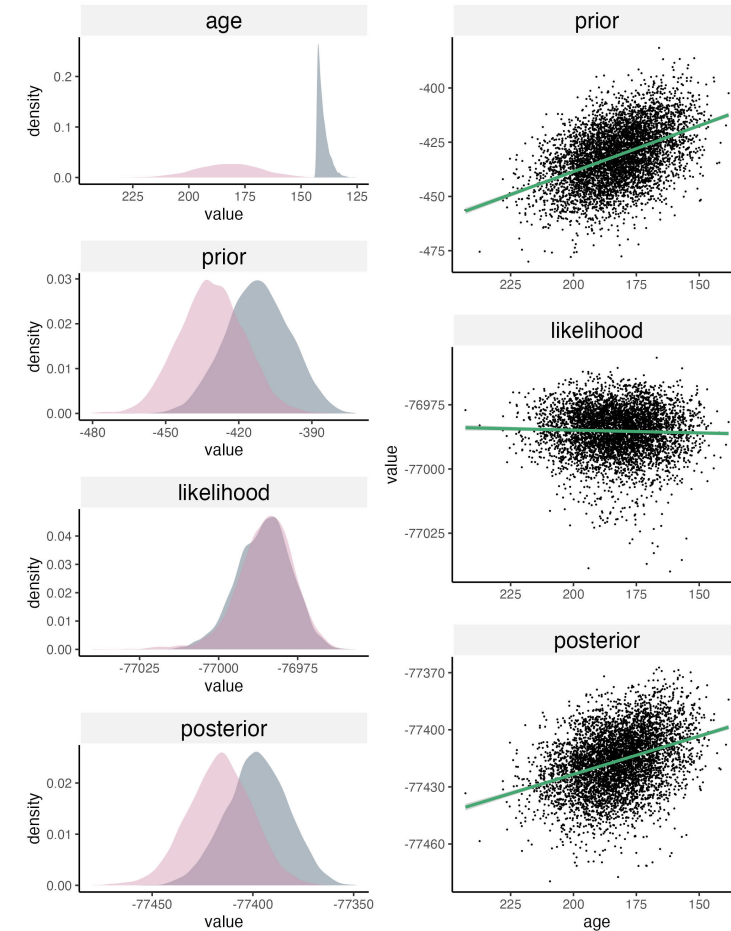
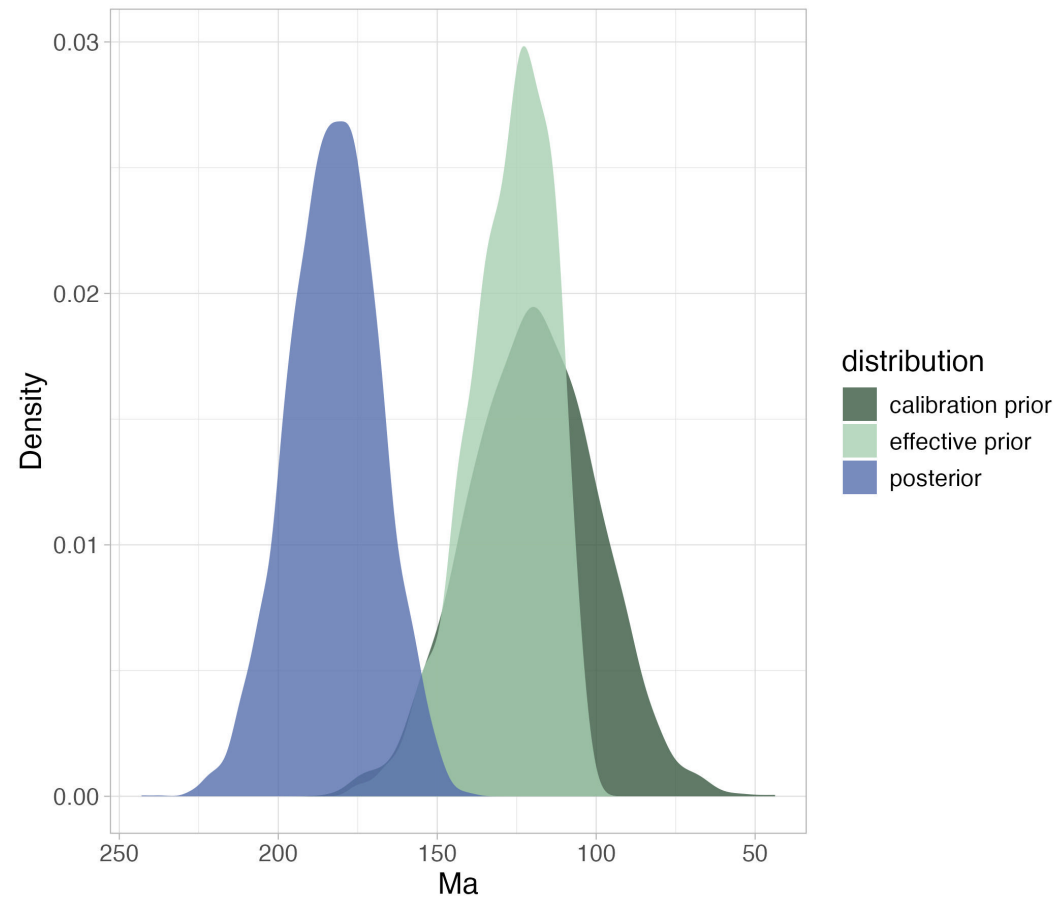
Article

Ancient Rhamnaceae flowers impute an origin for flowering plants exceeding 250-million-years ago

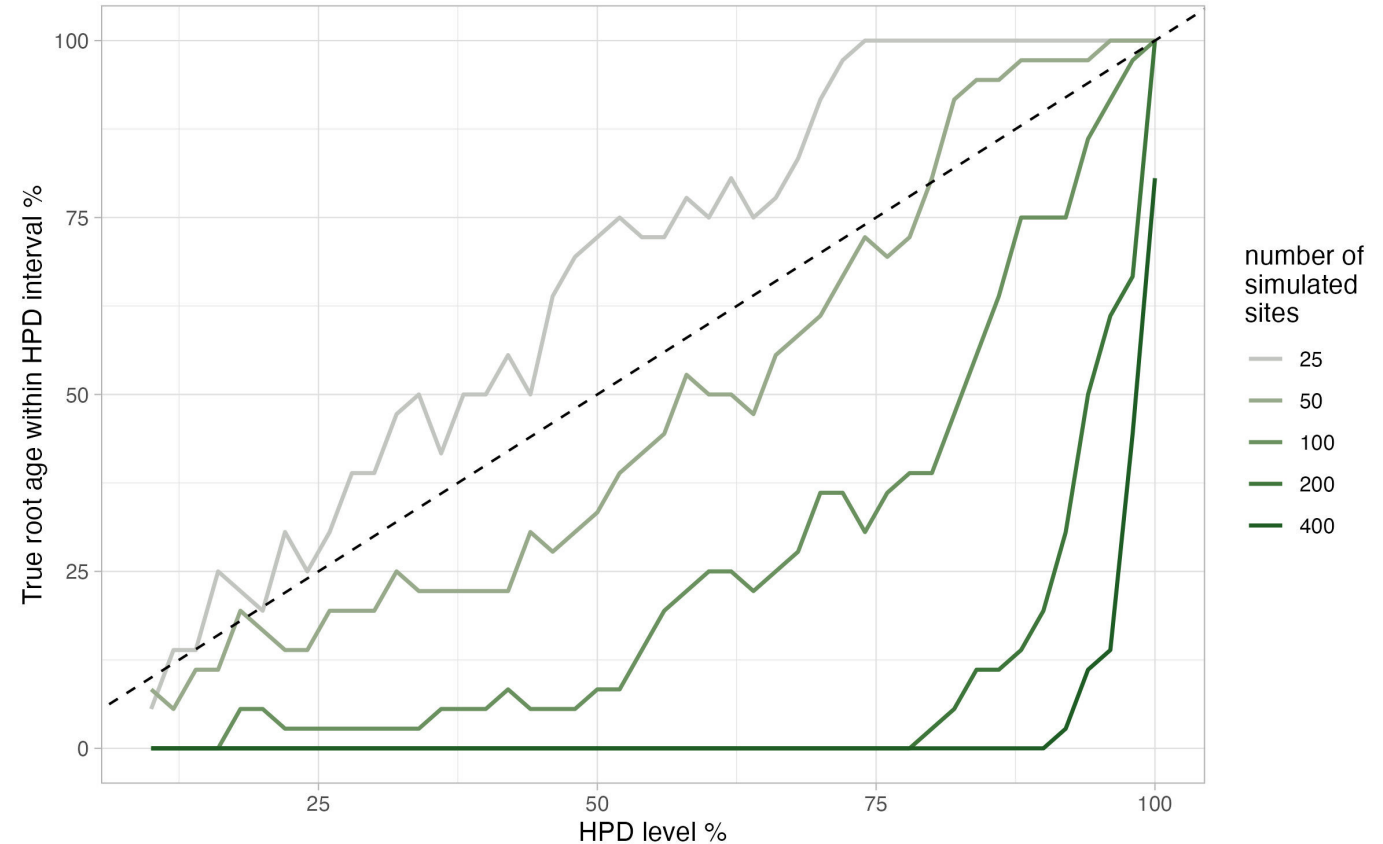
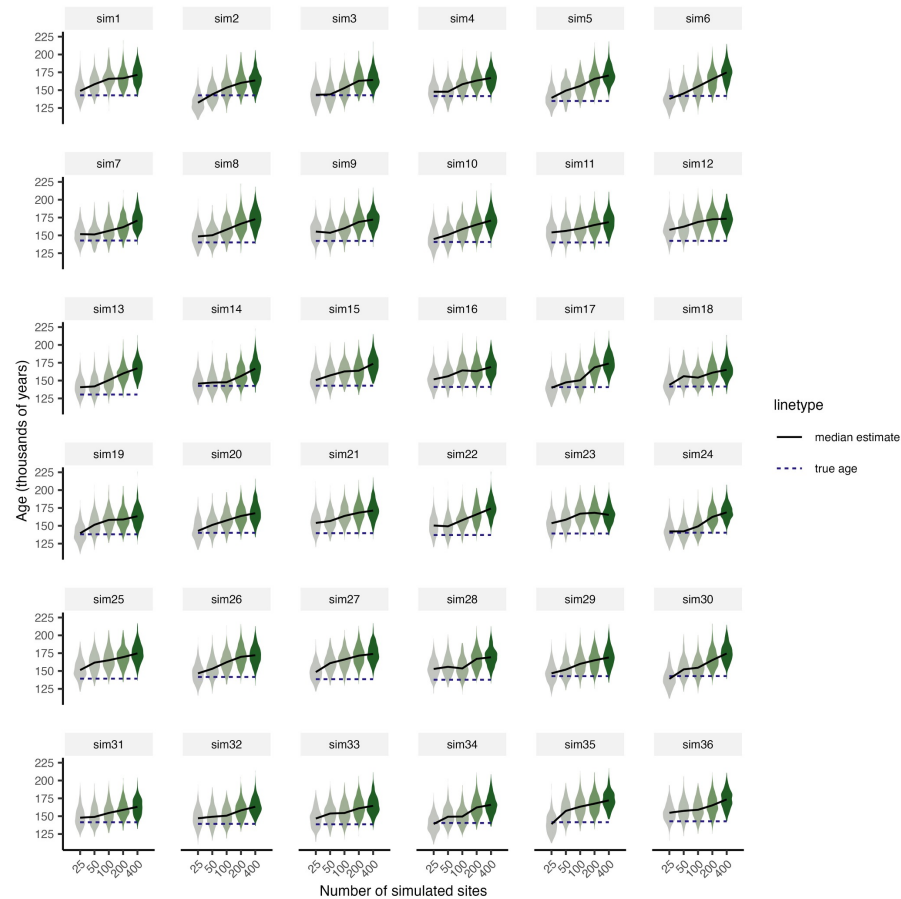
He and Lamont 2022



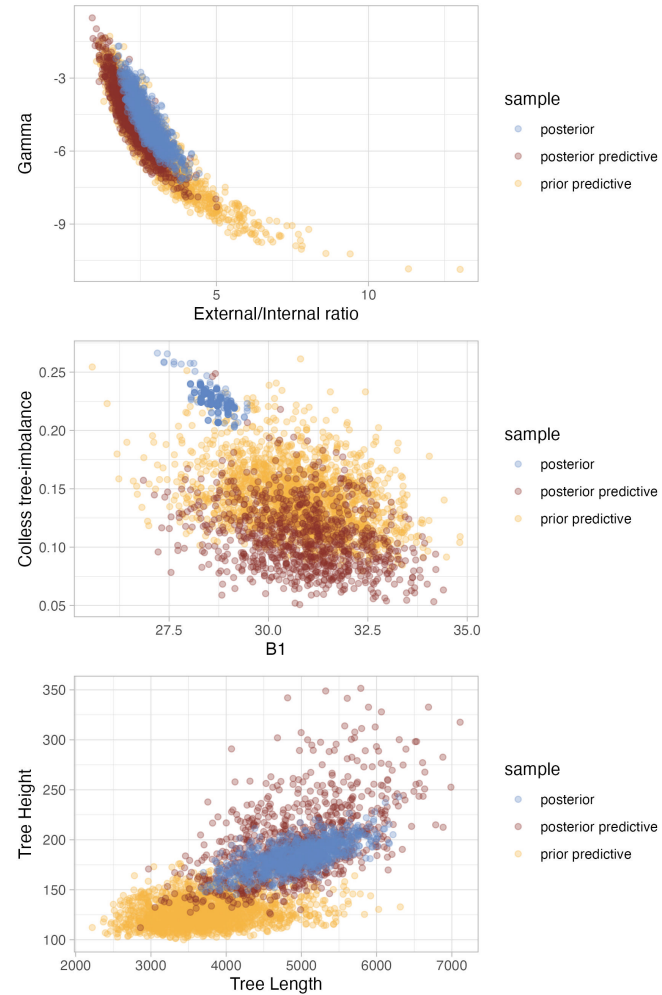
Ancient dates and flat likelihood, again



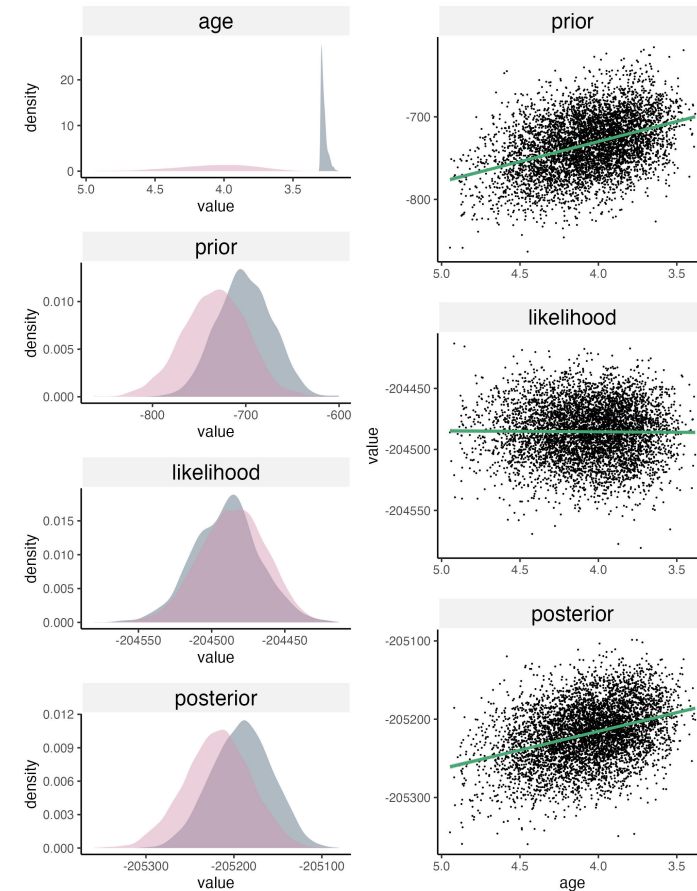
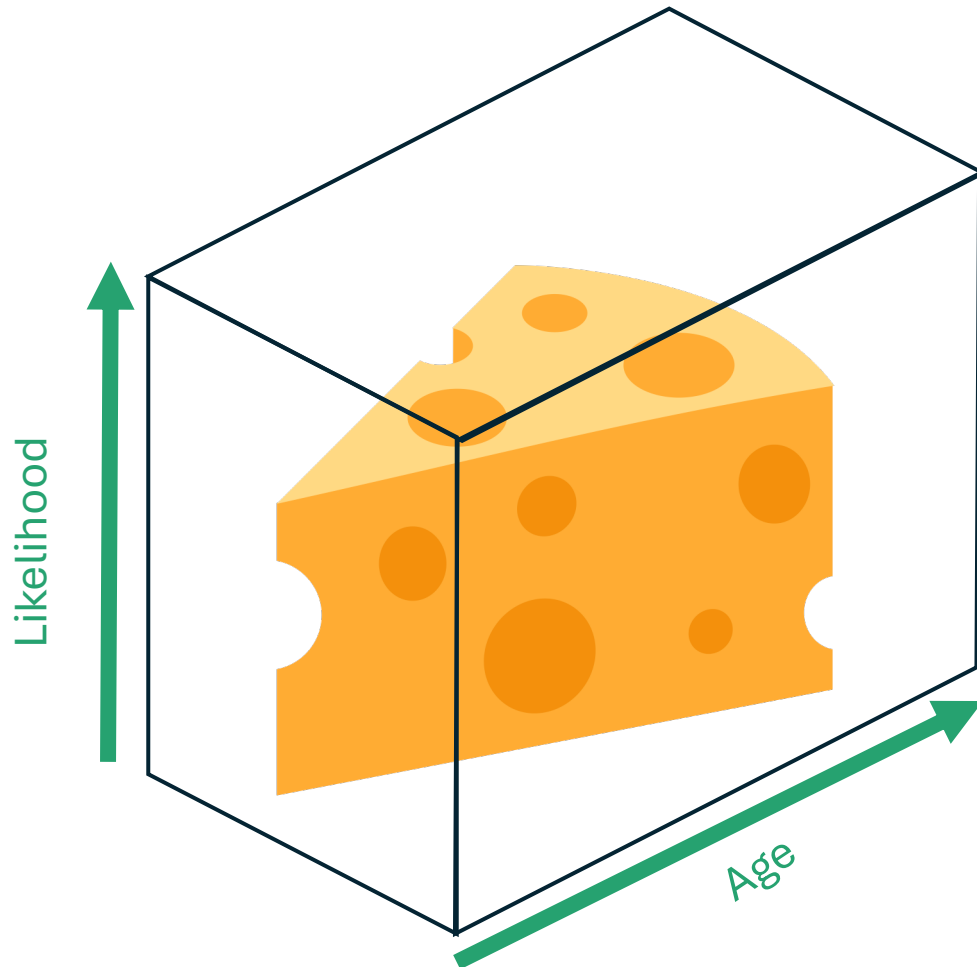
Simulations on posterior trees



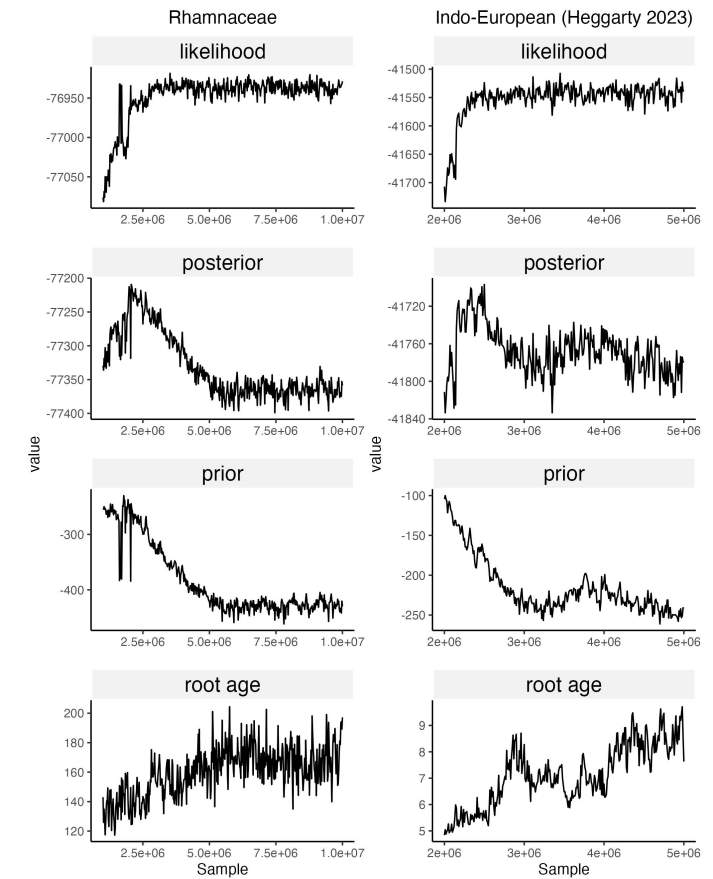
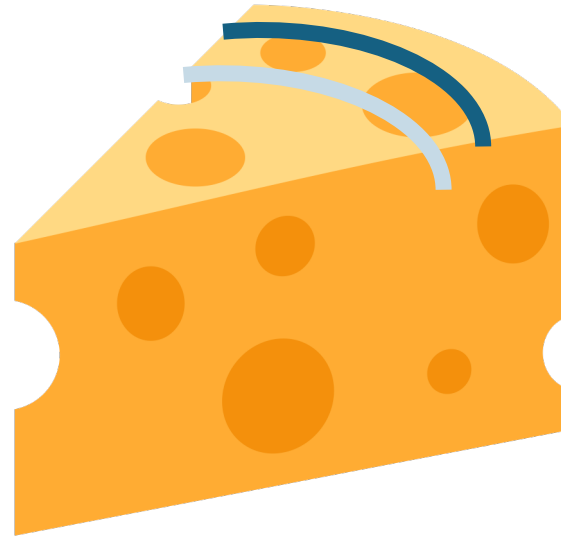
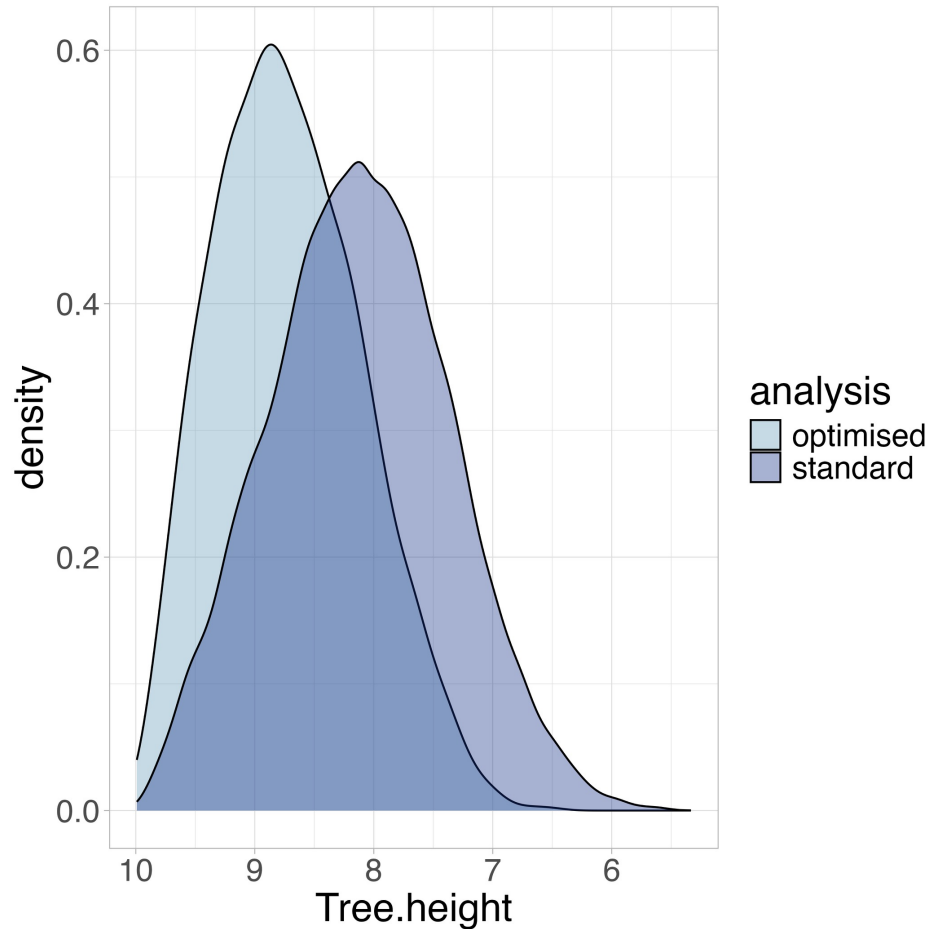
Rhamnaceae posterior predictive



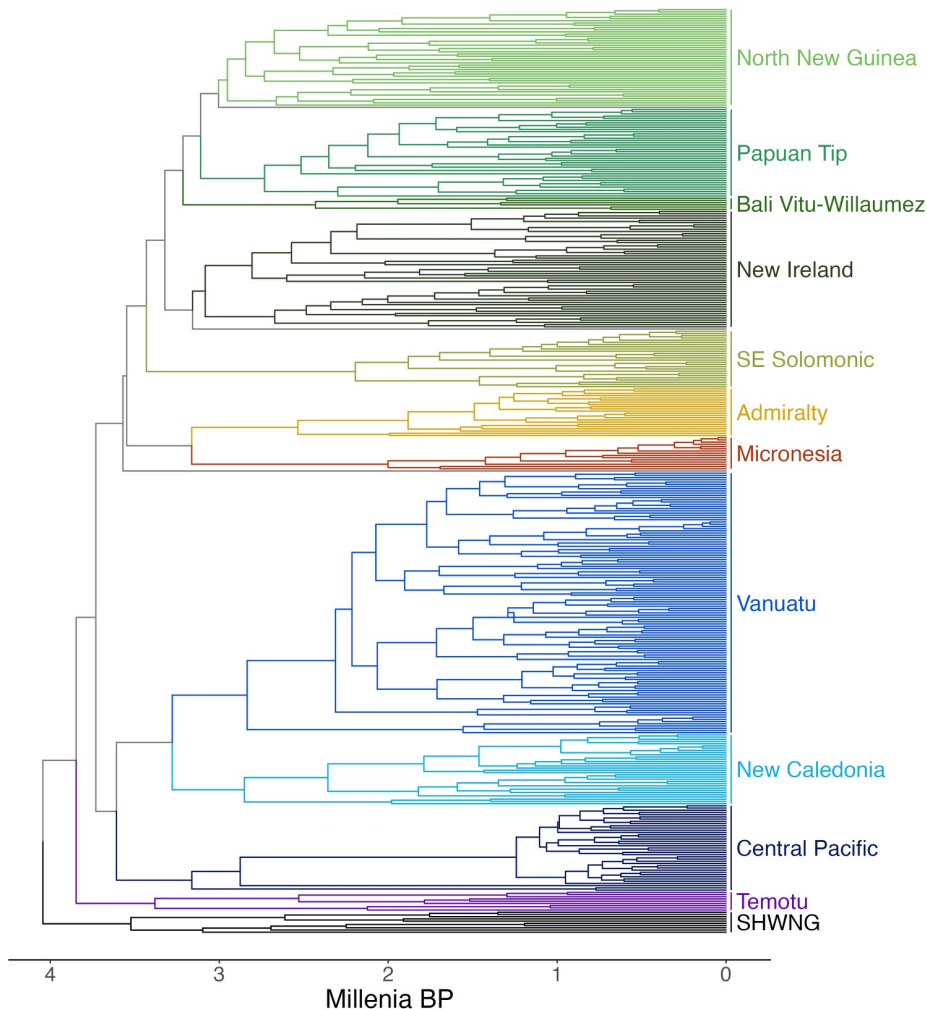
The expanding likelihood wedge



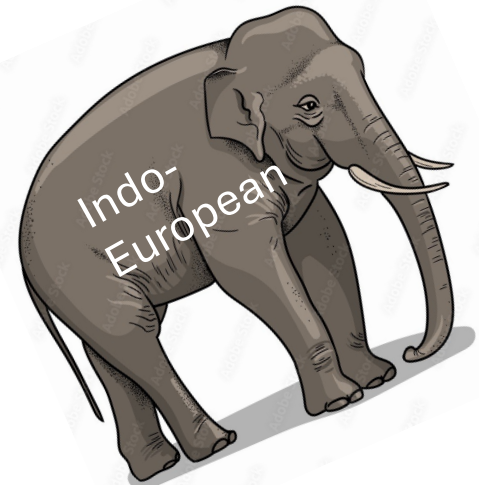
Sensitivity to software version



What can we say about Oceanic



- Results so far suggest that root age could be over-estimated
- Cannot draw conclusions about the early migration
- Alternative tree prior (CLADS) has not been successful



Conclusions

- Model mis-specification of tree priors could render molecular clocks uninformative
- This occurs across different calibration set-ups (tip and node) and data type (linguistic, DNA)
- Not clear how widespread

