

The Copenhagen phylolinguistics workshop

Workshop description

Phylogenetic inference is one of the central endeavors of historical linguistics. Research in this area is in the midst of a methodological revolution, which has been brought about by an influx of computational methods originally developed for the study of molecular evolution and the tree of life. Foremost among these new methods is Bayesian phylogenetic inference, which above all enables historical linguists to pursue questions that were previously intractable. This workshop provides a hands-on introduction to Bayesian phylogenetics with the statistical programming environment RevBayes. Although familiarity with linguistic change and the broad outlines of the history of Indo-European will be assumed, prior experience with Bayesian methods and programming will not.

Website

<https://rootsofeurope.ku.dk/kalender/arrangementer-2021/the-copenhagen-phylolinguistics-workshop/>

Dates and times

Location: Zoom

Dates: 3, 5, 7 May 2021

Time: 6.30–8.30 pm CEST / 9.30–11.30 am PDT

Software

Please make sure to download and install the following software before the course begins. The first three programs can be downloaded by clicking on their names.

- [RevBayes v1.1.1](#)
- [Tracer](#)
- [Figtree](#)
- A text editor such as Sublime Text, NotePad++, Atom, BBEdit, vim, or emacs.

Contact

For questions, please contact Thomas Olander (olander@hum.ku.dk) or David Goldstein (dgoldstein@humnet.ucla.edu).