

Rousing the BEAST: A molecular phylogenetic pipeline tutorial



Software Requirements:

Text editor:

- **Windows:** <https://notepad-plus-plus.org>
- **Mac:** <http://www.barebones.com/products/bbedit/>

Programs:

- Mesquite: <https://www.mesquiteproject.org>
- PAUP*: <https://paup.phylosolutions.com>
- BEAST2: <http://www.beast2.org>
- TRACER: <http://tree.bio.ed.ac.uk/software/tracer/>
- FigTree: <http://tree.bio.ed.ac.uk/software/figtree/>

Also, be sure you **make an account** on the CIPRES portal (<https://www.phylo.org>).

Please download these programs *before* the tutorial. The directions for each are relatively straightforward; if you have any difficulties, please let me know at zhancock@bio.tamu.edu or (903) 809-3625.

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