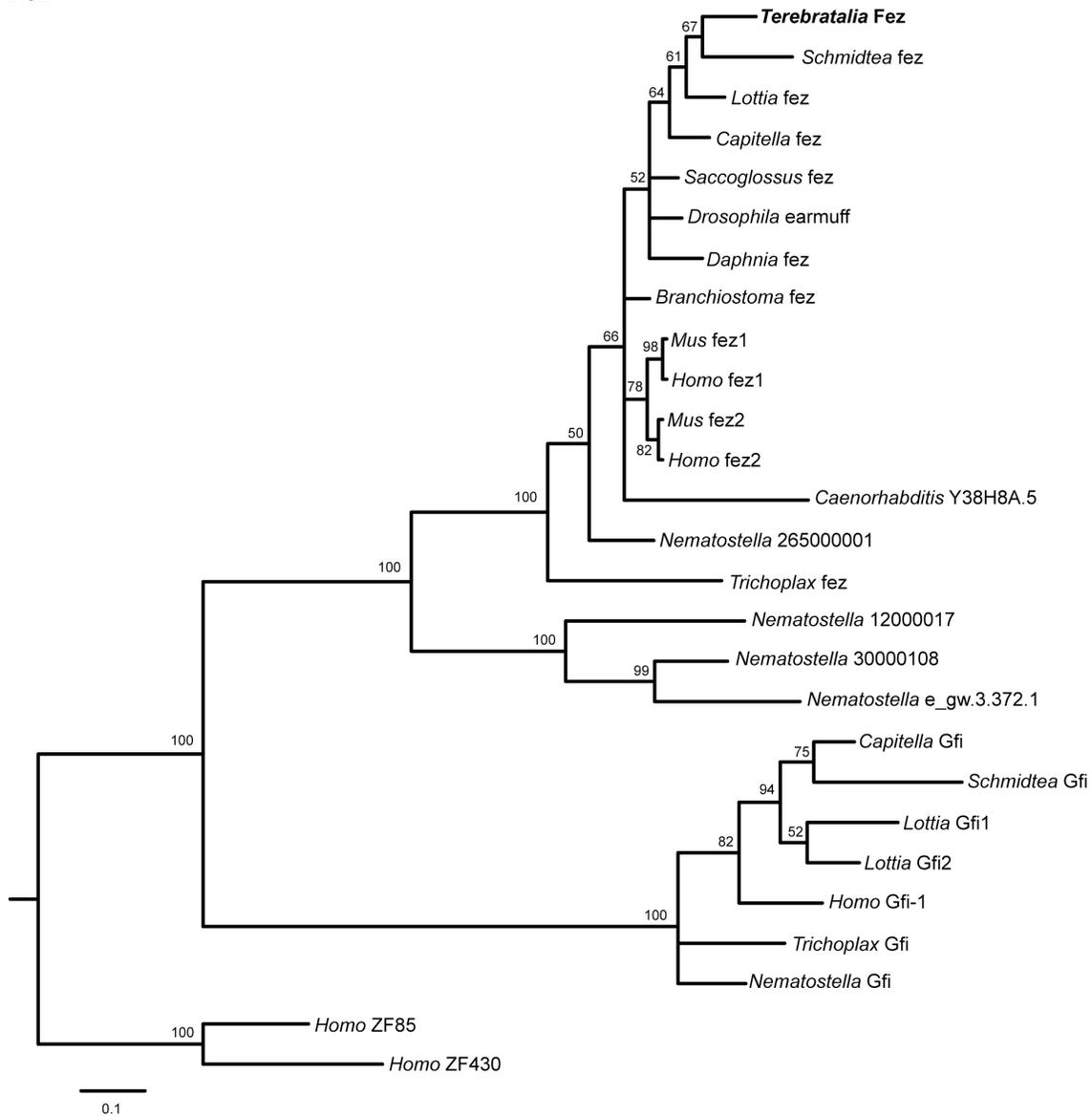


## Fez



### Additional File 3: Phylogenetic analysis of *Tt-fez*.

Phylogram of *Tt-fez* and related zinc-finger proteins, supporting the orthology assignment of *Tt-fez*. Posterior probability for the Fez clade, including *Tt-fez*, is 100 percent. The phylogram is a consensus of the last 2,000,000 generations from a Bayesian likelihood analysis with four independent runs of 10,000,000 generations each.