

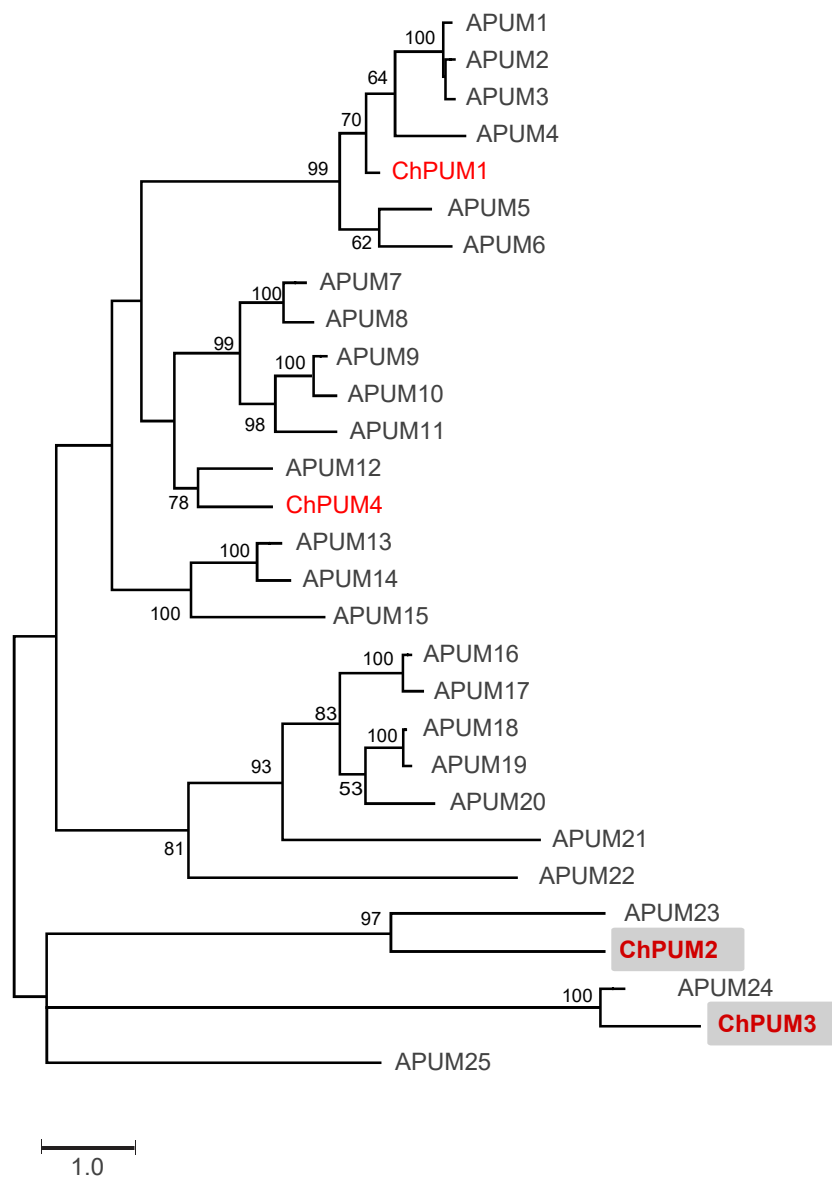


Figure S1. Phylogenetic tree of the Pumilio proteins using 25 APUMs from *A. thaliana* APUMs and 4 ChPUMs from *C. corallina*. The maximum likelihood tree was generated using the JTT+F+G model with 1000 bootstrapping replicates.