



A maximum-likelihood tree (model GTR + I + G; $\ln L = -21068.86248$; Best of NNIs and SPRs) of three-finger toxins is presented here. Node support was evaluated through 1000 bootstrap replicates. Branches with node support of more than 850/1000 replicates are indicated as thick lines. Site-model 8 computed omega and the total number of positively selected sites detected by its Bayes Empirical Bayes (BEB) approach ($PP \geq 0.95$) are also indicated for each toxin clade.