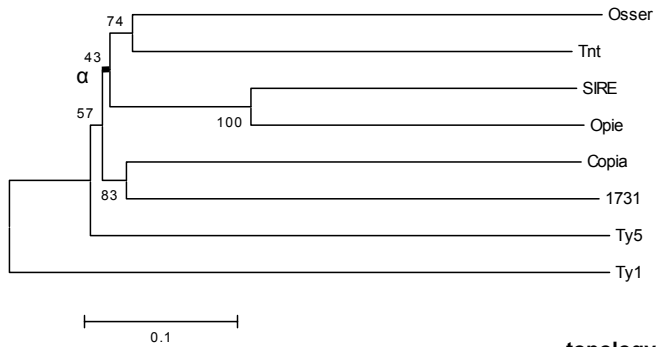
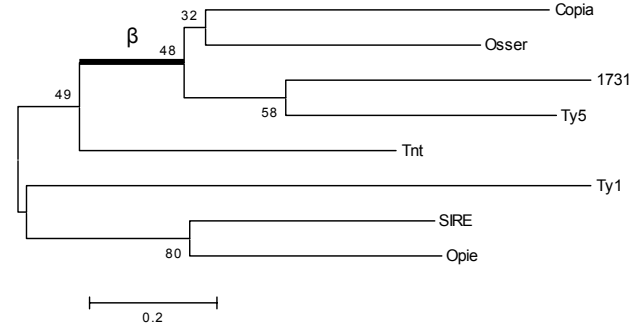


A Phylogenetic tree from full length sequence alignment



B Split cluster from sequence alignment in window (166-215)



C

