

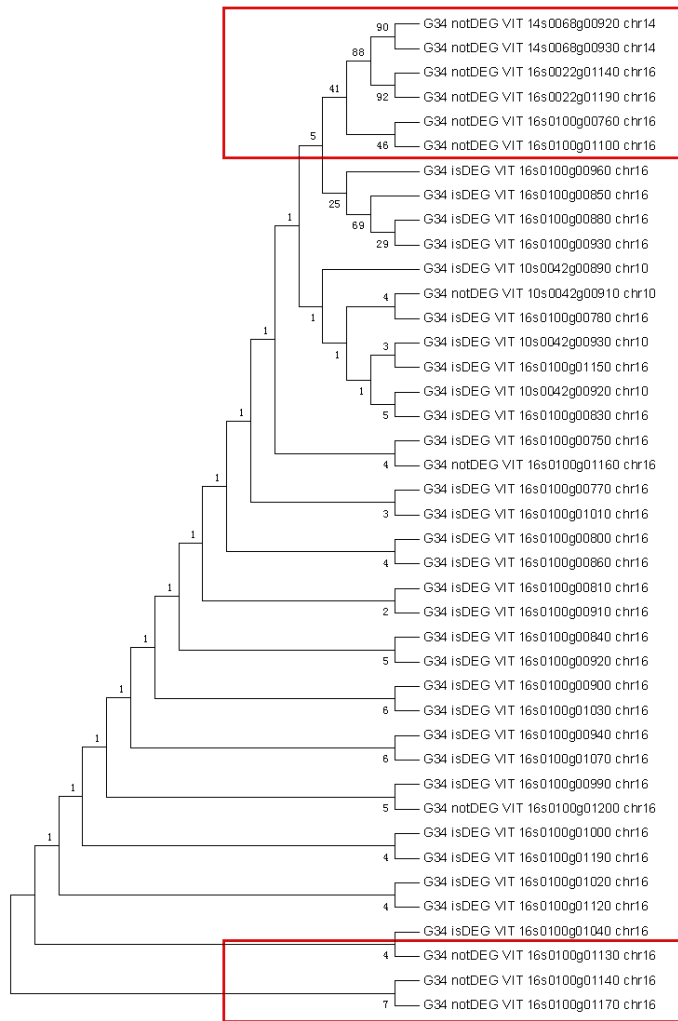
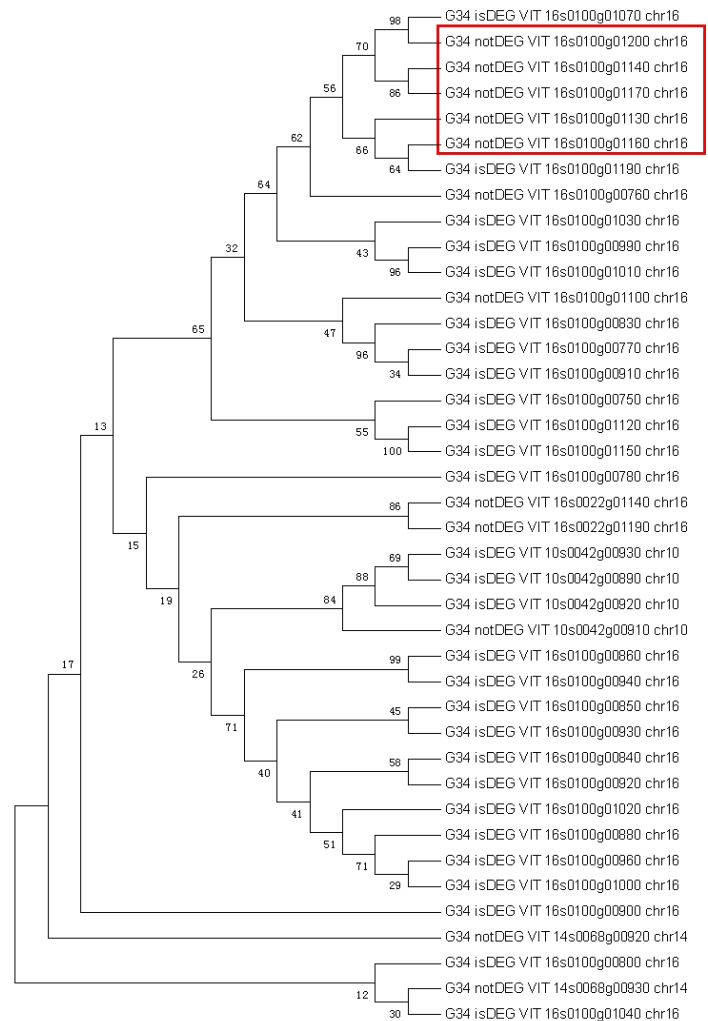
A**B**

Figure S20. The maximum-likelihood trees of the protein sequence (A) and the 1000 nt upstream of the transcription start sites (B) of 42 tandem duplicated STS genes in gene clusters of G34 families.