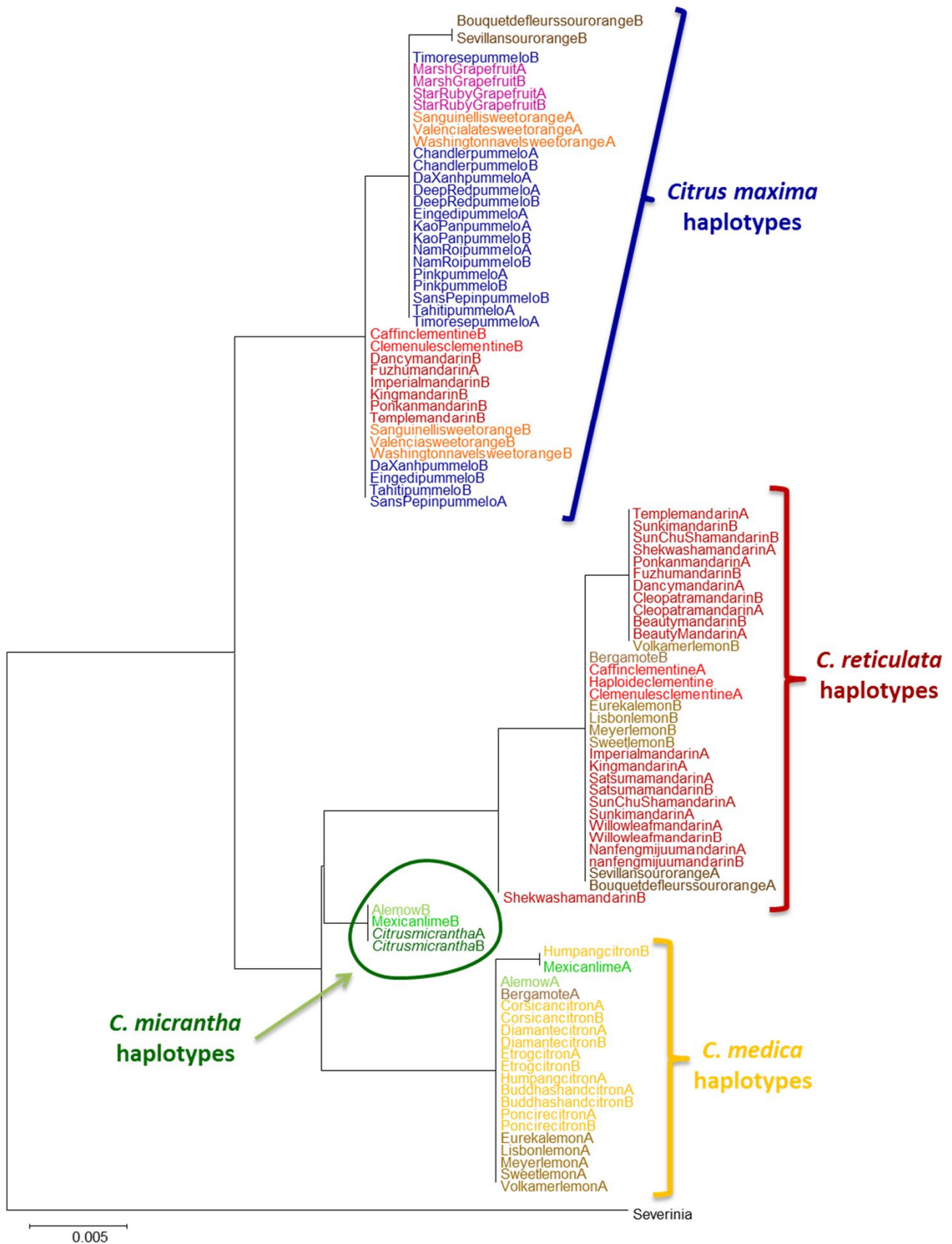


Next generation haplotyping to decipher the nuclear genomic interspecific admixture in Citrus species; analysis of the Chromosome 2. Curk et al. BMC genomic, Additional file 10.



Additional File 10: Maximum likelihood phylogenetic tree of the haplotypic data of the 2P35391362 gene fragment.