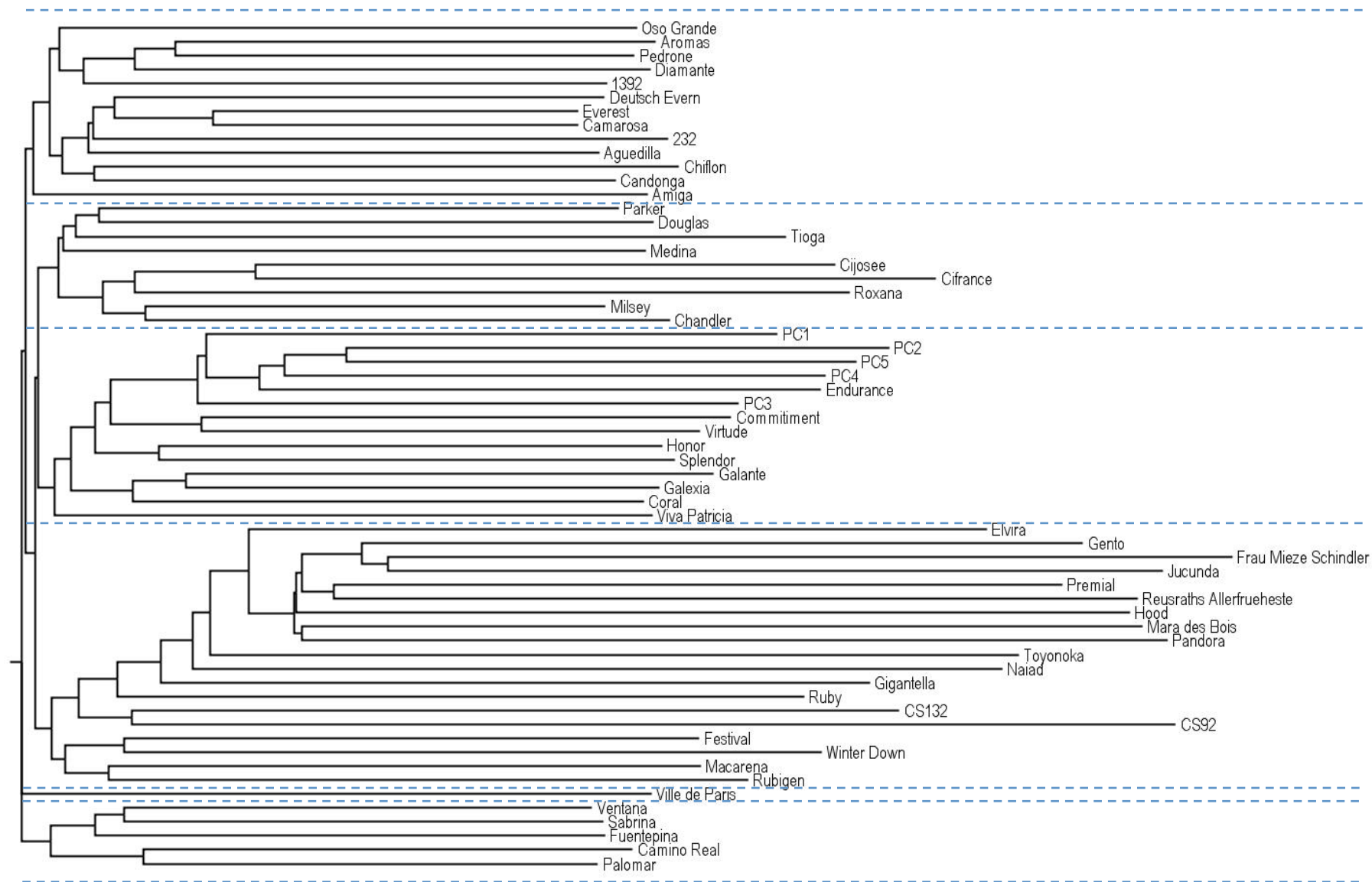




Online Resource 1 Neighbor-Joining clustering of 61 strawberry accessions (out of the 71 used in this manuscript) based on 21,971 SNPs markers. Accessions were selected based in an ongoing study (unpublished results) using a total of 176 accessions genotyped using the Affymetrix IStraw90 Axiom array (Bassil and Davis et al. 2015). The 61 selected accessions were also distributed in different clusters in the complete analysis.