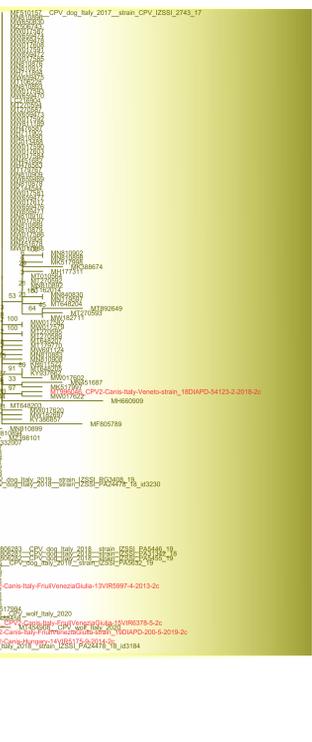
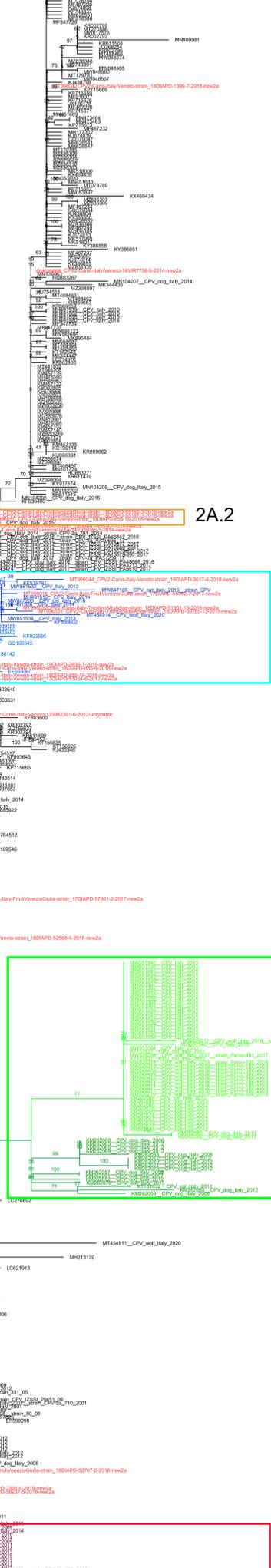


2C.2



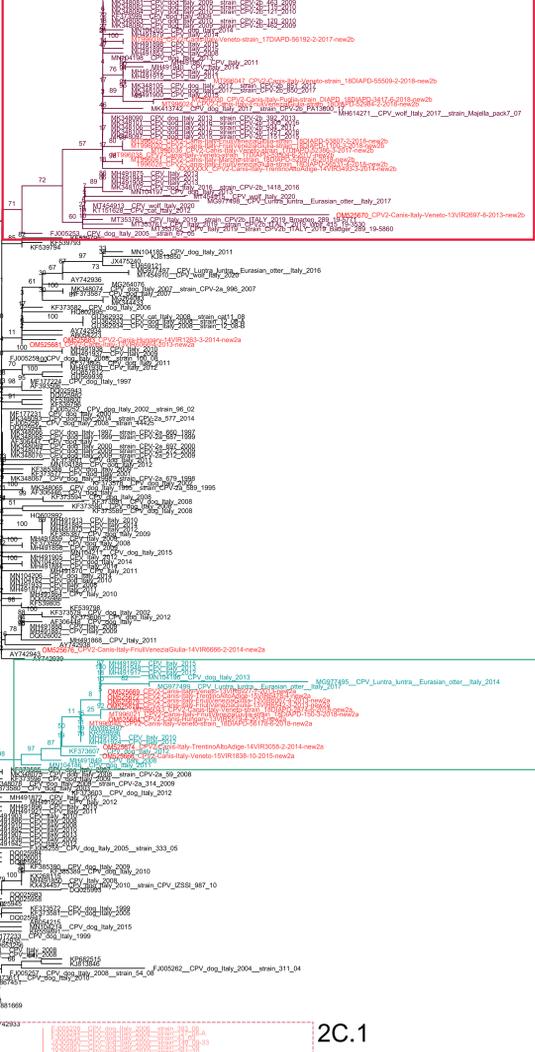
2A.2

2A.3

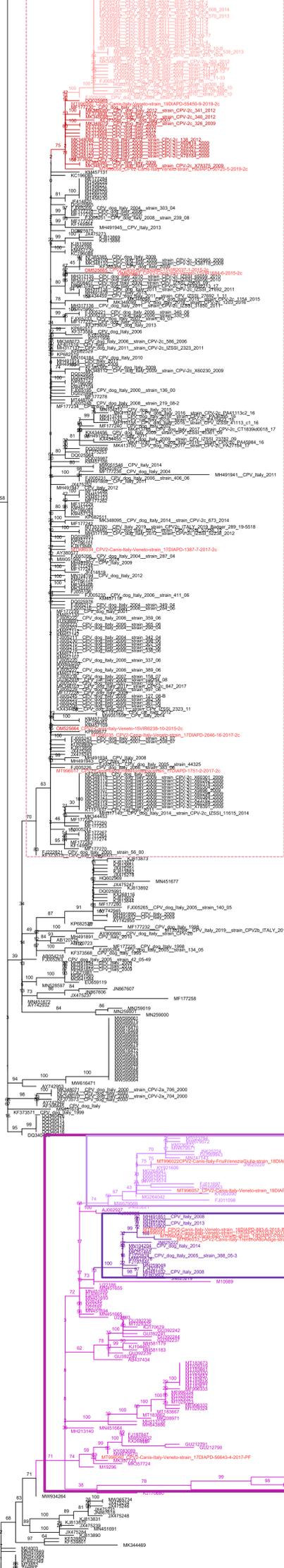


2B

2A.1



2C.1



vaccinal strains

OR

INT

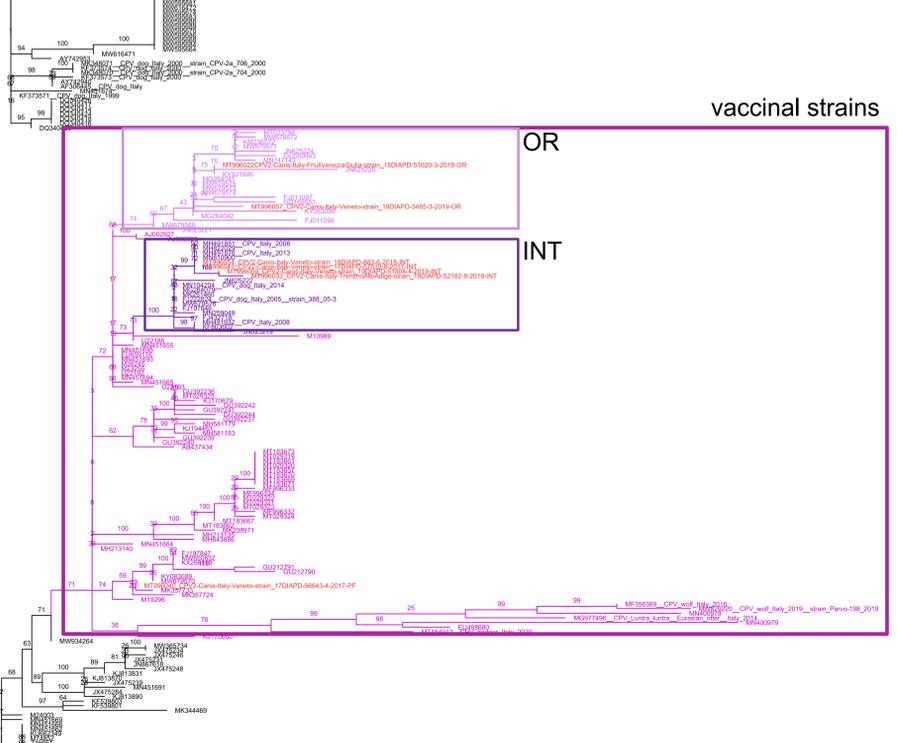


Figure S1. Phylogenetic tree of the VP2 nucleotide region of CPV-2 identified in the present study (highlighted in red) and similar CPV-2 strains available in GenBank. In rectangular boxes are reported eight monophyletic groups containing the majority of the IZSve CPV-2 strains.