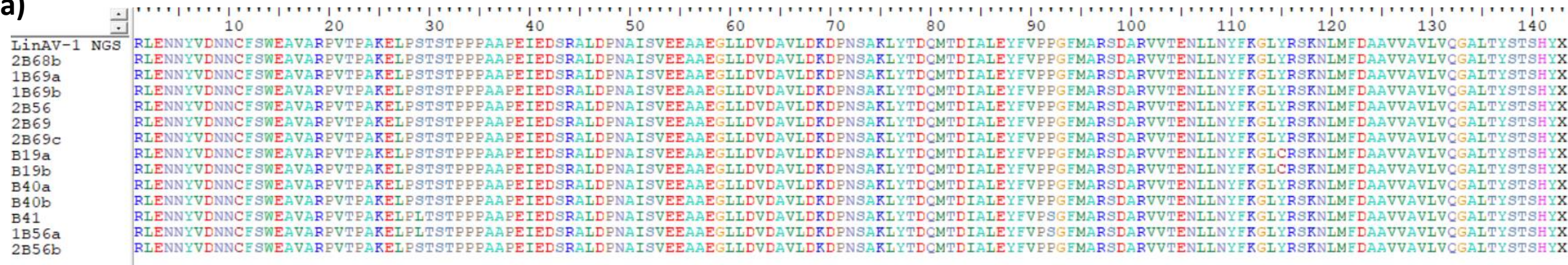


(a)



(b)

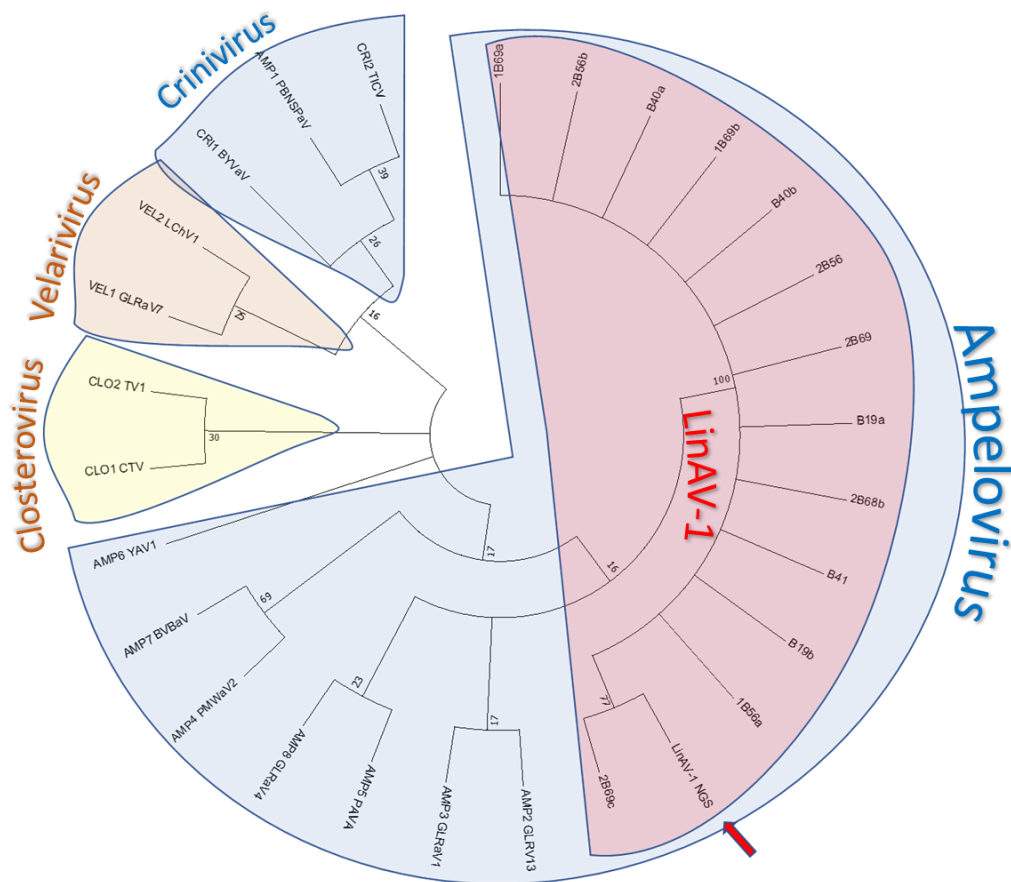


Figure S2: Validation of the detection of LinAV-1 in plants collected along Cantillana preserve by the alignment (a) and the phylogenetic tree constructed by maximum parsimony (b) of the deduced amino acid sequences of a partial region of CP gene.