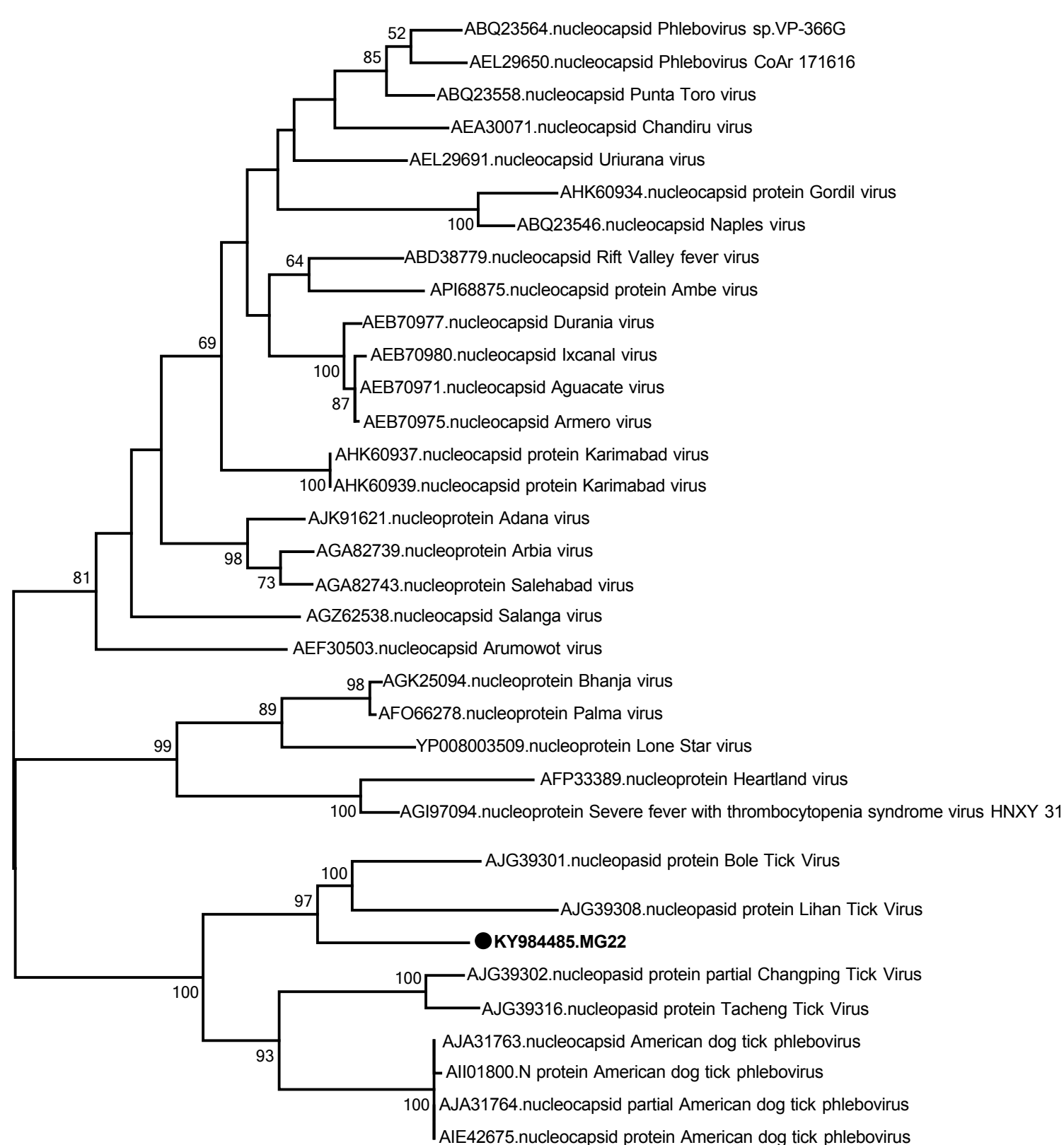




Additional file 3: Figure S1. The maximum likelihood analysis of the partial putative nucleocapsid protein (152 amino acids) of phleboviruses. The sequence characterized in this study is given in bold and indicated with the GenBank accession number, pool code and a black circle. Global virus strains are indicated by GenBank accession numbers, virus and strain/isolate names. Bootstrap values higher than 60 are provided.