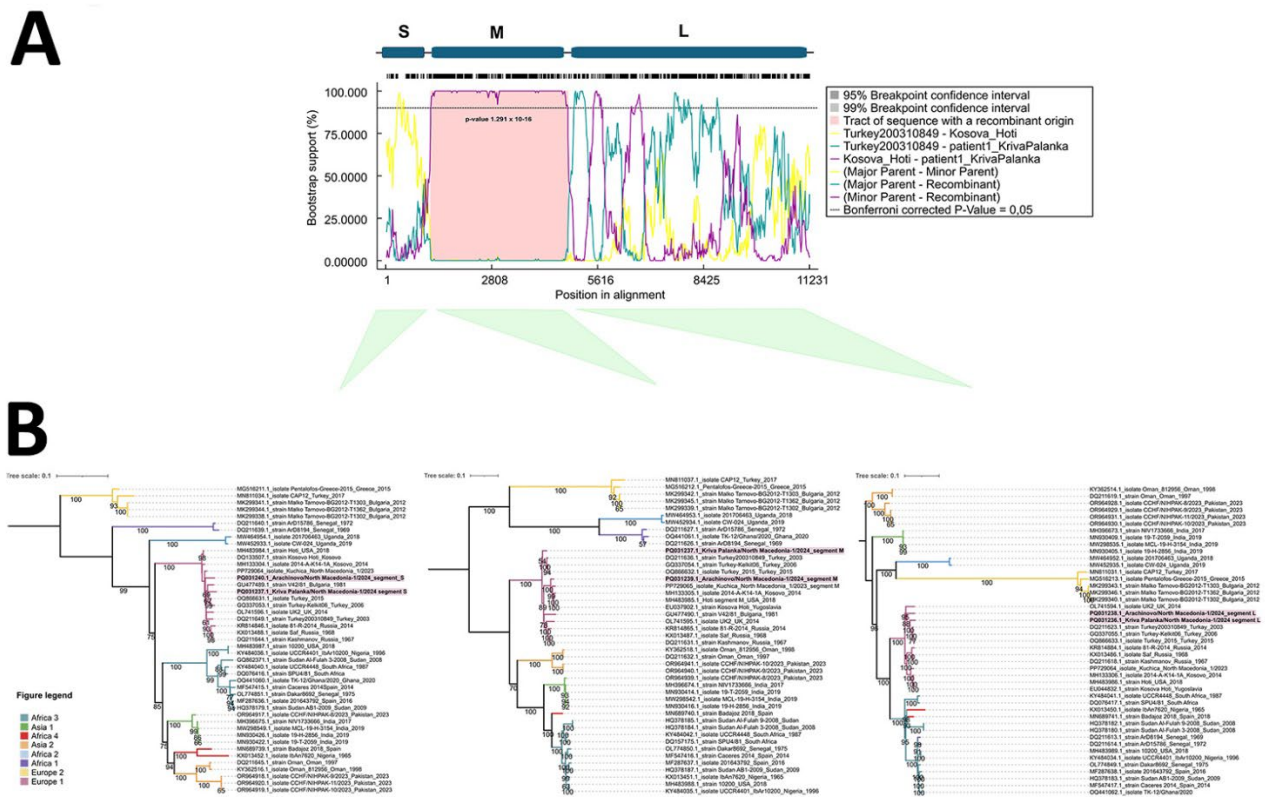


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# Two Human Infections with Diverse European Crimean-Congo Hemorrhagic Fever Virus Strains, North Macedonia, 2024

## Appendix 2



**Appendix 2 Figure.** Bootsacan and phylogenetic analysis of Crimean–Congo hemorrhagic fever virus isolates, North Macedonia, April–May 2024. A) Bootsacan analysis of the CCHFV isolate Arachinovo/North Macedonia-1/2024 (GenBank accession nos. PQ031238.1–PQ031240.1) and Kriva Palanka/North Macedonia-1/2024 (GenBank accession nos. PQ031235.1–PQ031237.1). Approximate positions of the genes are also depicted above the Bootsacan plot. B) Maximum-likelihood phylogenetic trees of the 3 viral genomic segments. Sequences examined in this study are highlighted with colored background. L, large; M, medium; S, small.