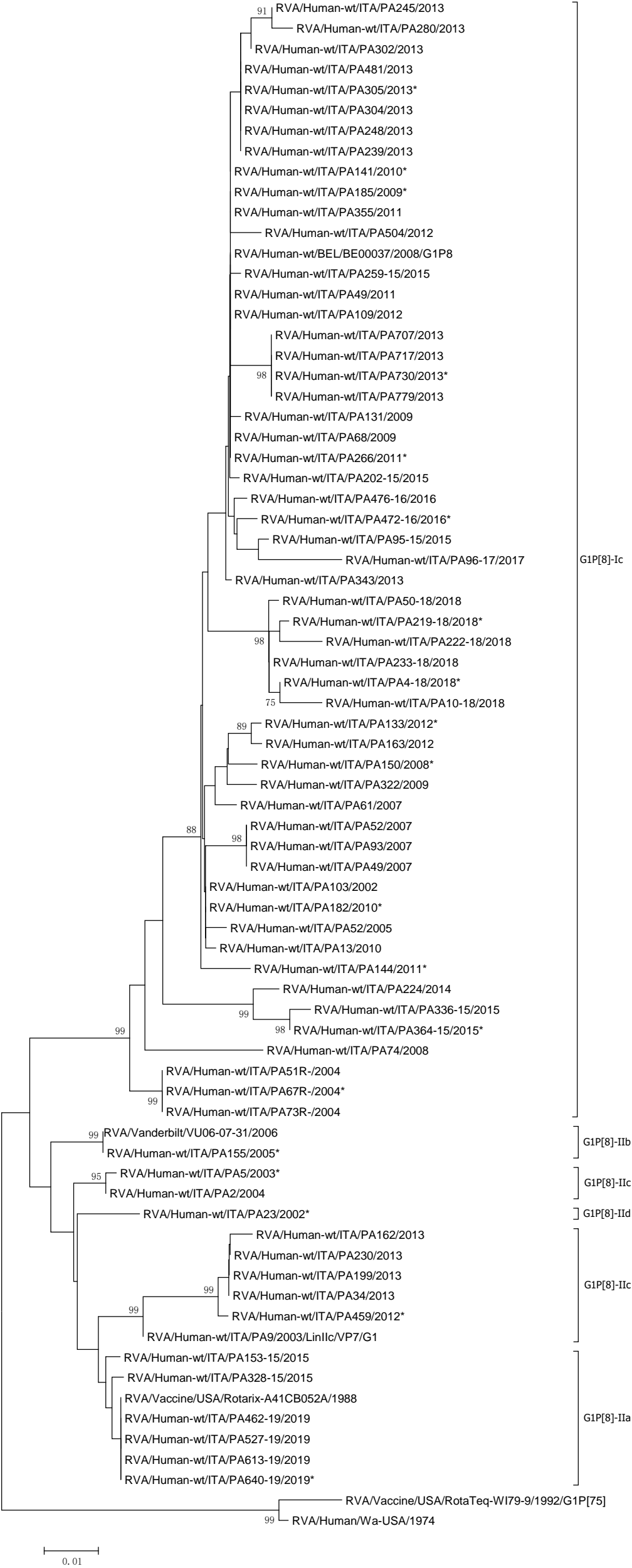
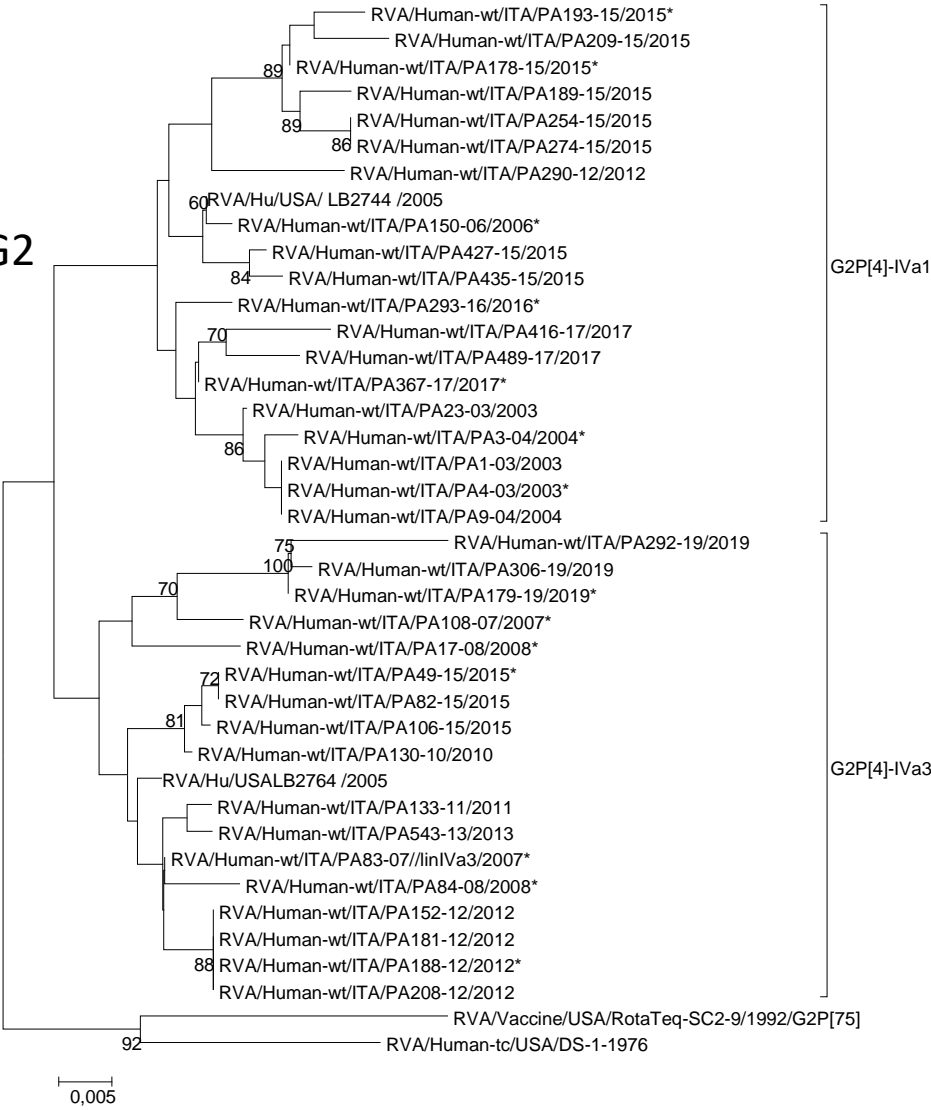


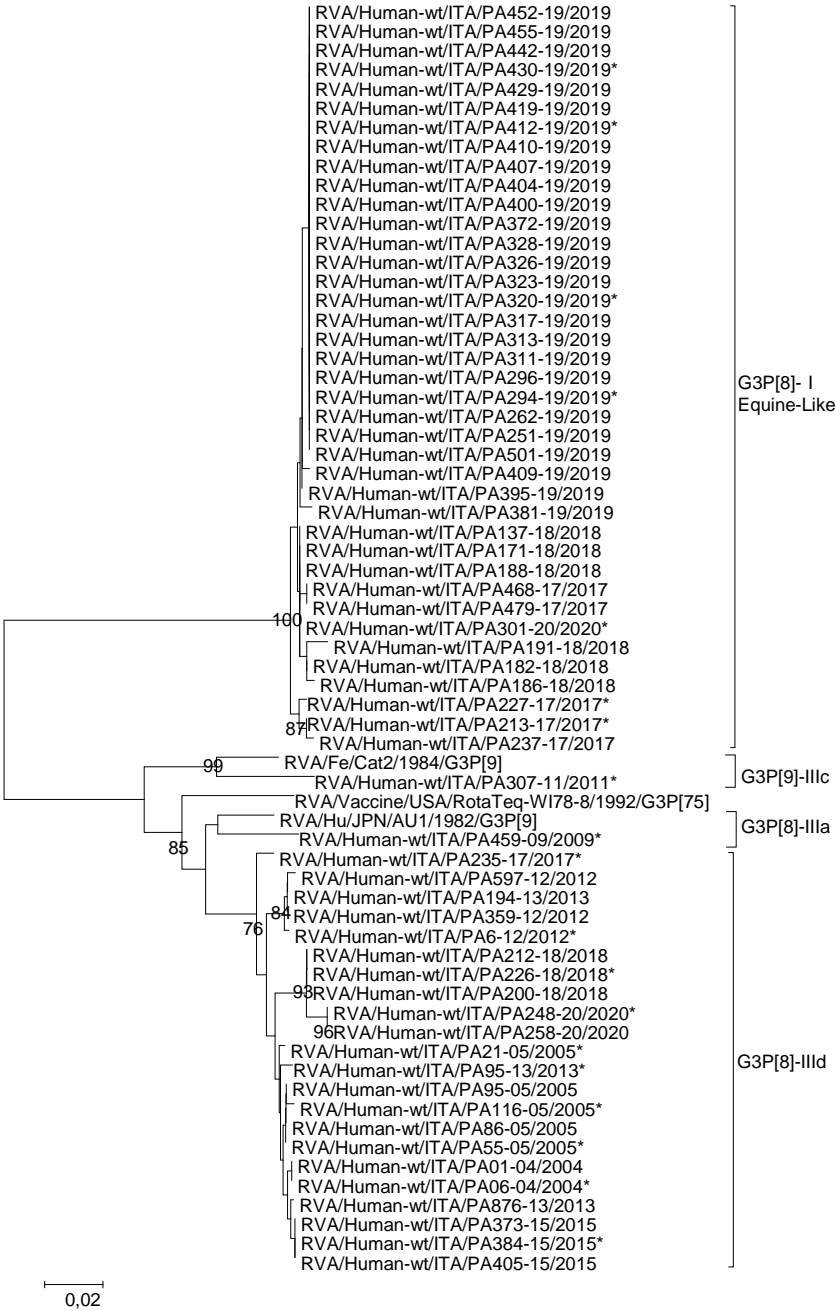
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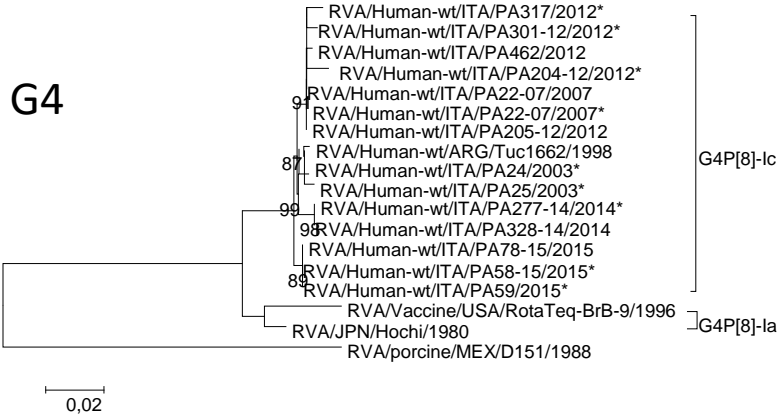
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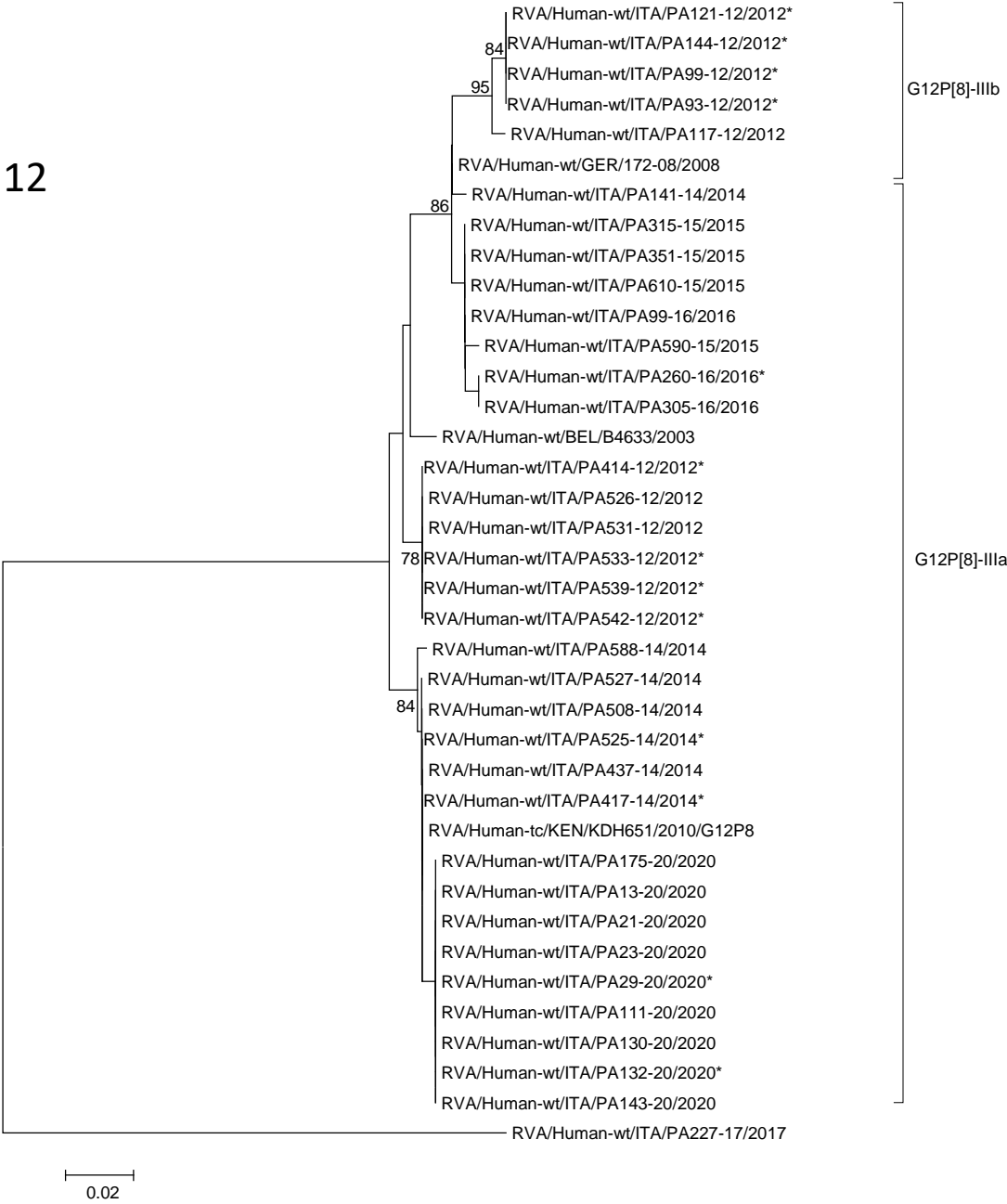
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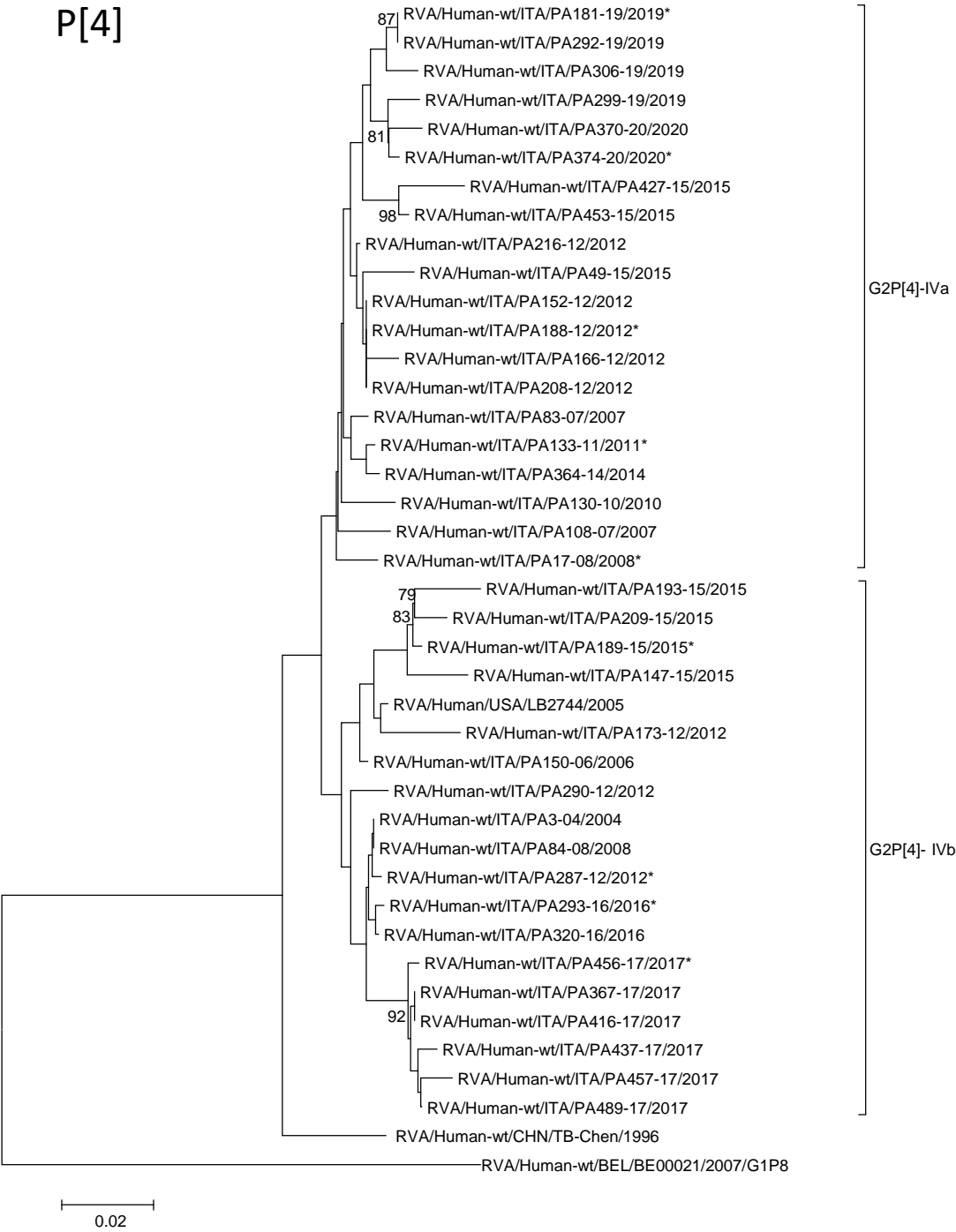
G9



G12



P[4]



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G1P[8]-III

Figure S1: Phylogenetic analysis of partial VP7 (G1–4, G9 and G12) and VP4 (P[4]–P[8]) nucleotide sequences of Italian strains detected in the study. The phylogenetic trees were built using the neighbor-joining method and Kimura's two-parameter model, and bootstrapped with 1000 repetitions. Bootstrap values 75% are indicated. * Sequences availables in GenBank, under the following accession numbers (VP7, G1:ON525269 - ON525287, G2:ON525288 - ON525295, G3:ON525296 - ON525309, G4:ON525310 - ON525314, G9:ON525315 - ON525325, G12:ON525326 - ON525328; VP4, P4:ON525329 - ON525336, P8:ON525337 - ON525350)