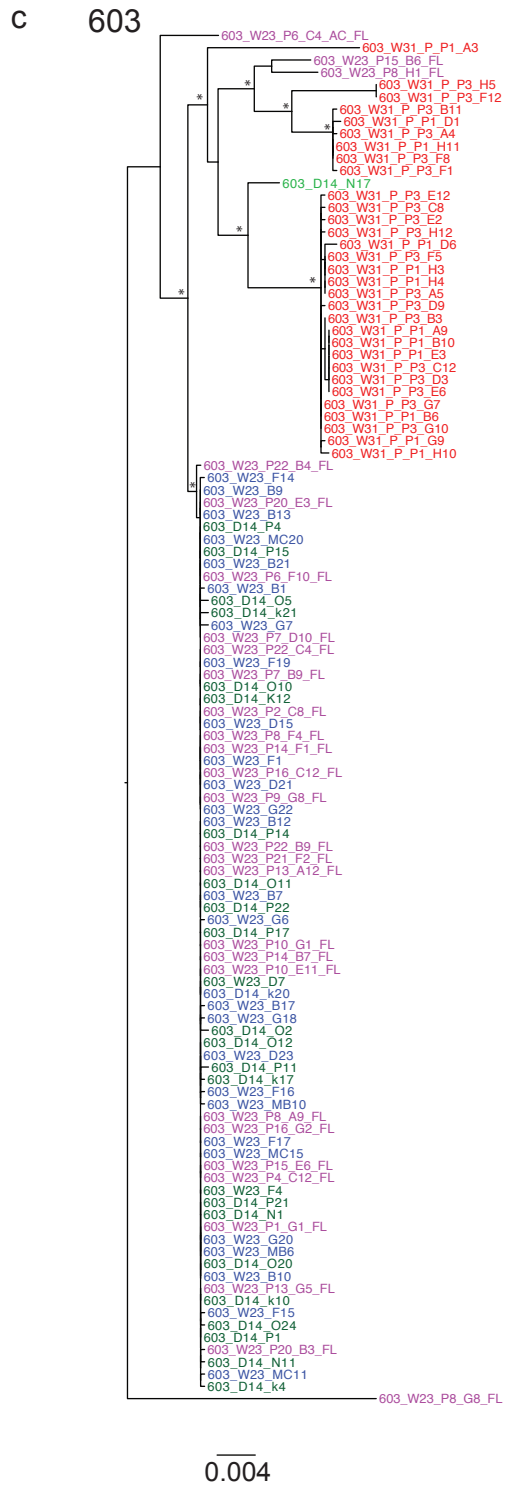
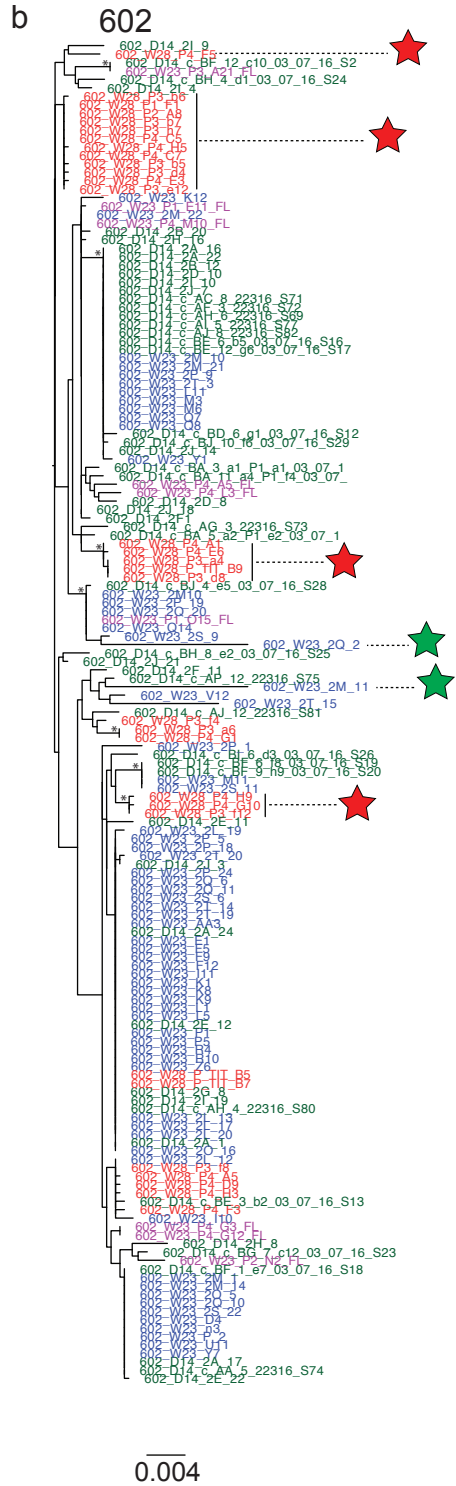
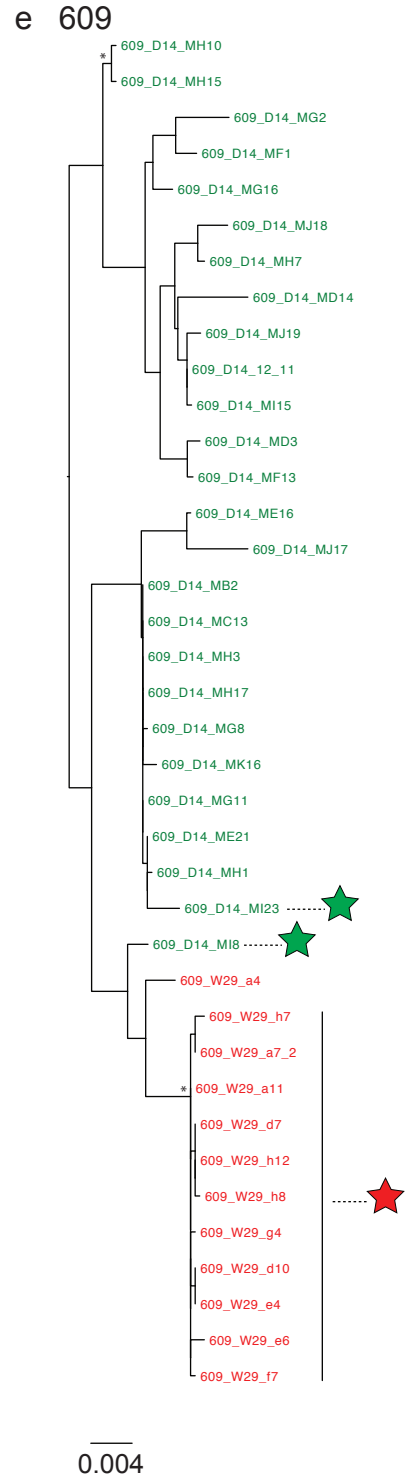
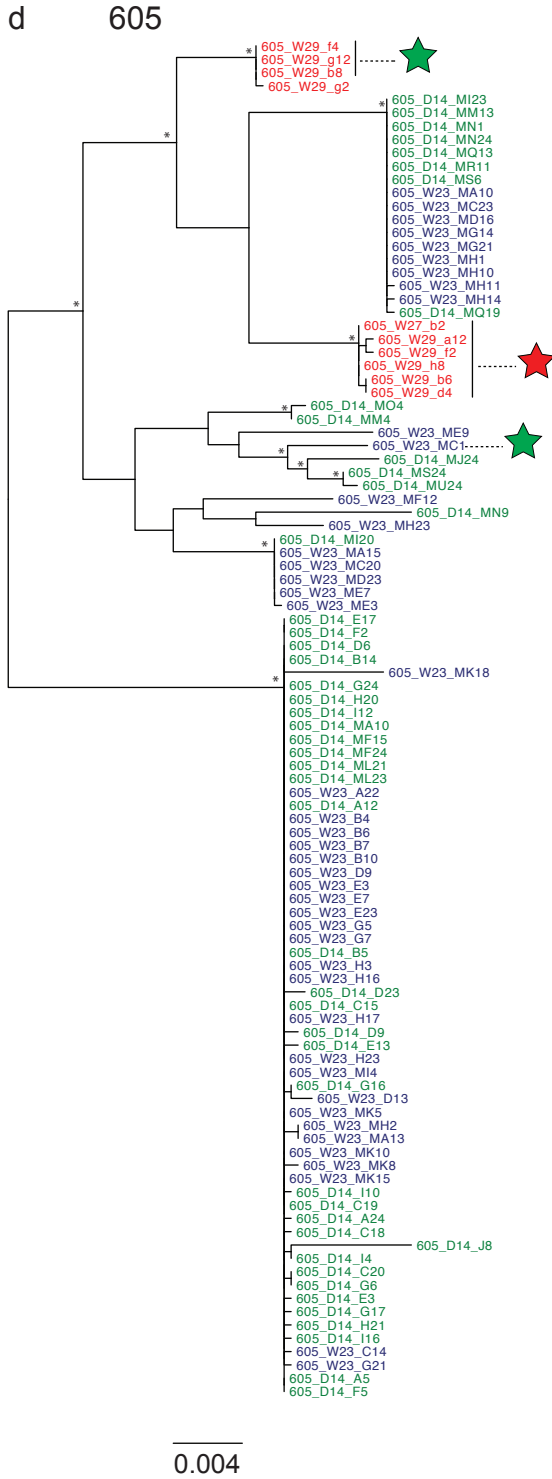


Supplementary Figure 7. Maximum likelihood phylogenetic trees of full-length *env* sequences from Q²VOA cultures, rebound SGA, and near full-length genome (NFL). Viruses from week -2 are represented in green, from week 23 in blue, rebound SGA in red, and from NFL in purple. Asterisks indicate nodes with significant bootstrap values (bootstrap support $\geq 90\%$). Green stars represent the recombinant parents and red stars represent the child recombinant for the examples shown in Fig. 5b.

Supplementary Figure 7 continued.

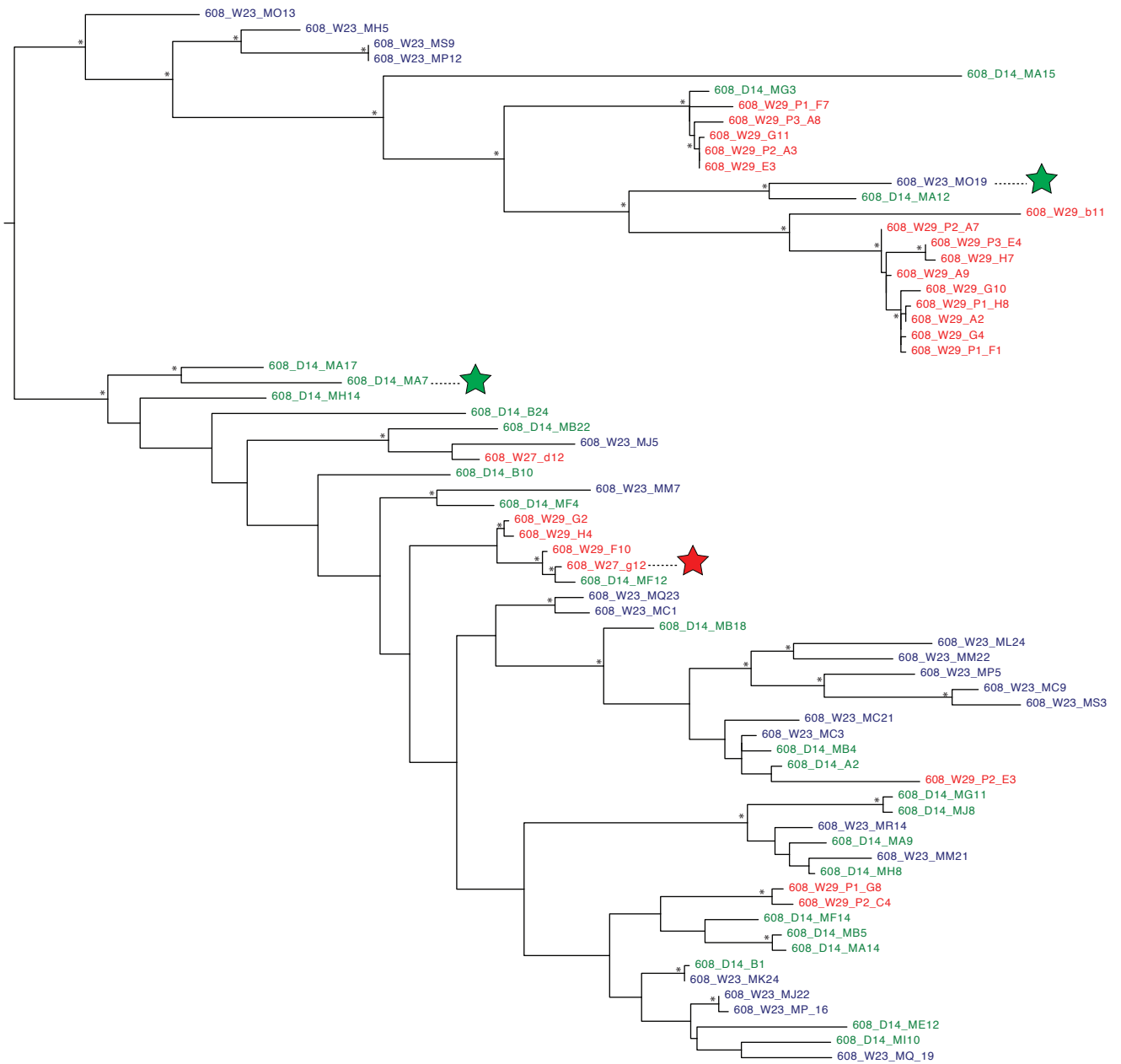


Supplementary Figure 7 continued.



Supplementary Figure 7 continued.

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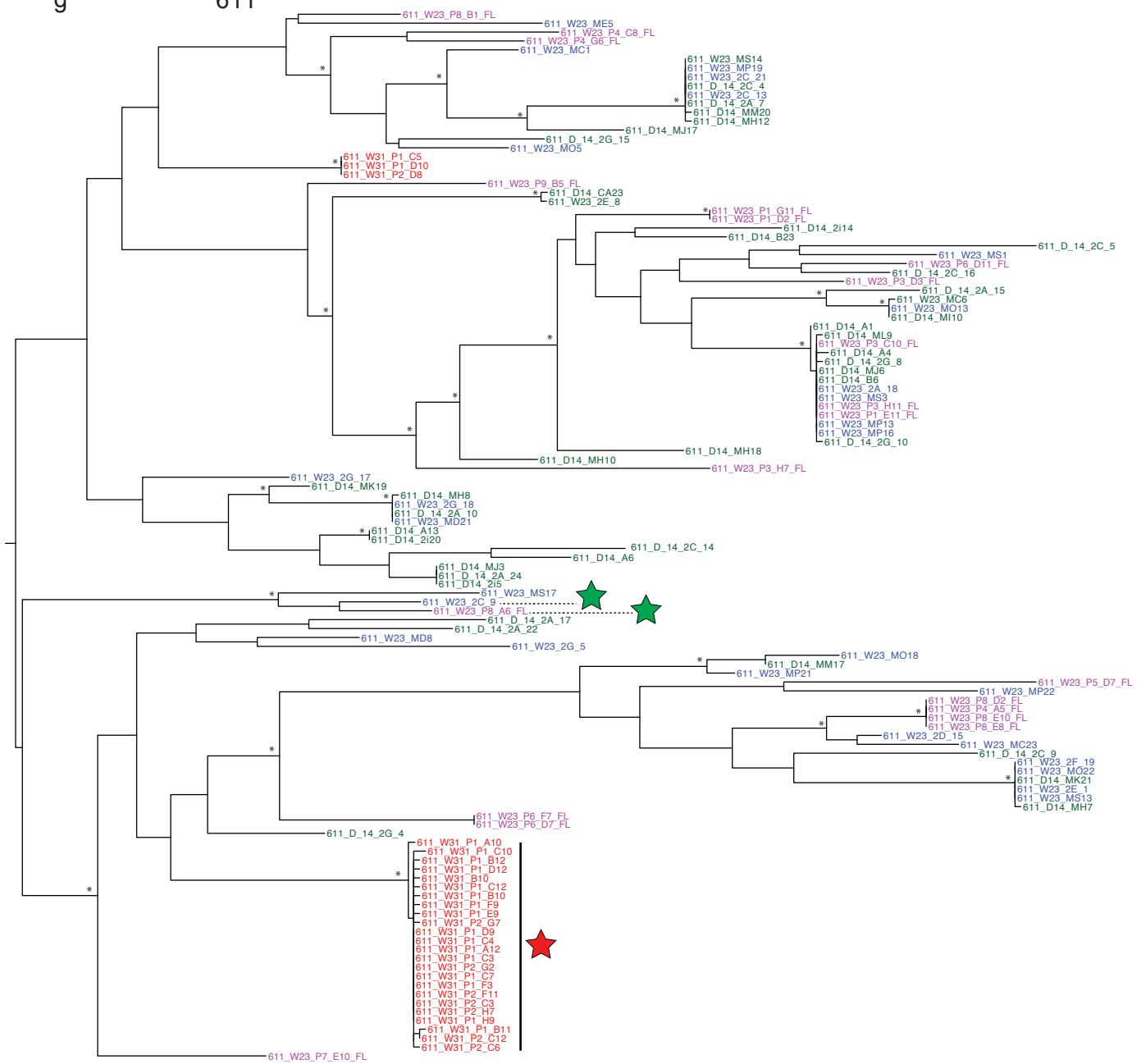


0.004

Supplementary Figure 7 continued.

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611



0.004

Supplementary Figure 7 continued.

