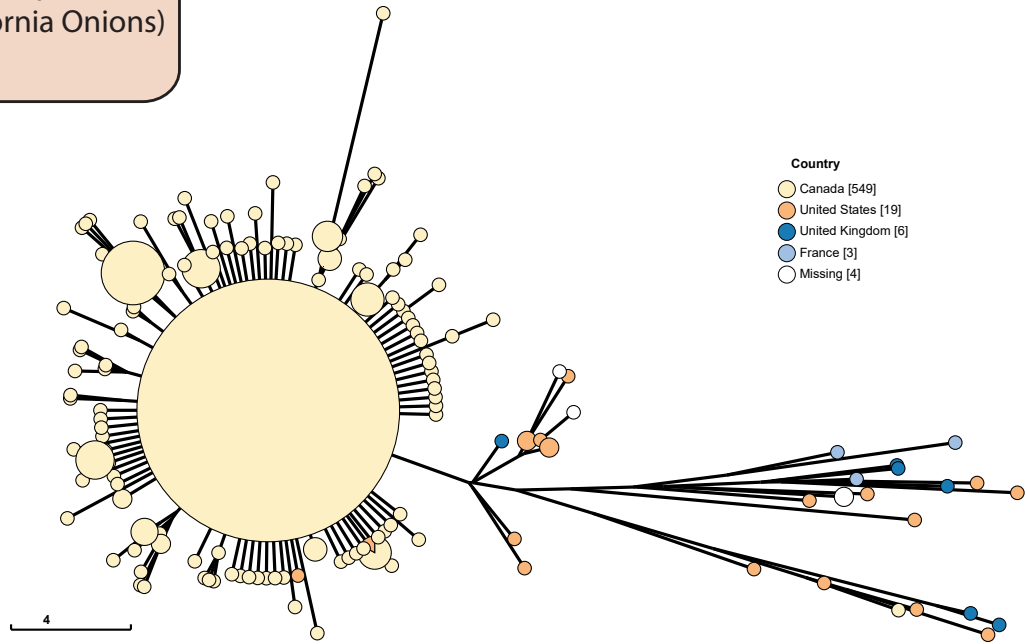


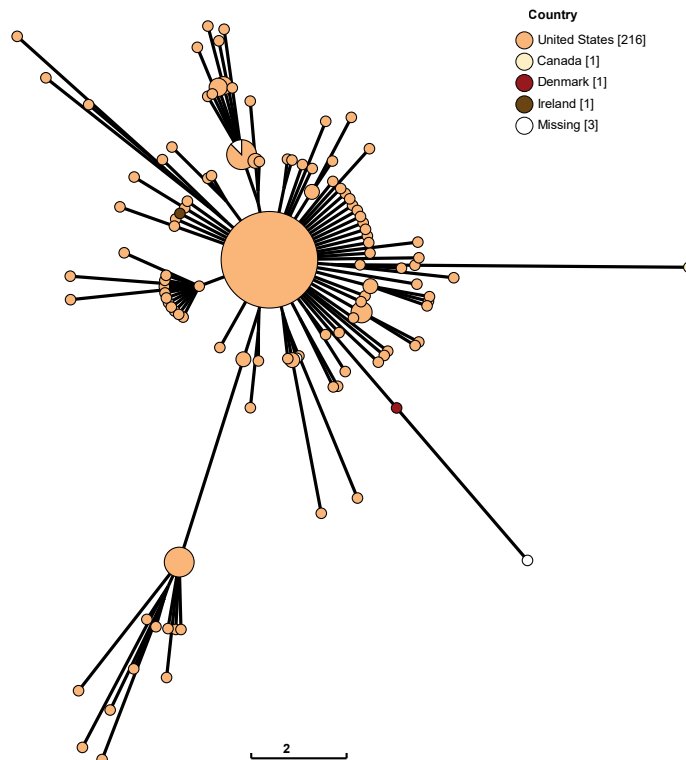
A

HC20_4179 Newport
Human (California Onions)
2000-2020



B

HC20_39803 Hadar
Human
2000-2018



S8 Fig. Ninja NJ visualization of allelic differences in the 3002 core genes of the cgMLST *Salmonella* scheme with GrapeTree for genomes within A) HC20_39803 and B) HC20_4179. Further information on these HC20 clusters is summarized in Tables 2, 4 and 5, and an interactive version of both trees can be accessed at https://enterobase.warwick.ac.uk/ms_tree?tree_id=45421 and https://enterobase.warwick.ac.uk/ms_tree?tree_id=60146.