



S7 Fig. Ninja NJ visualization of allelic differences in the 3002 core genes of the cgMLST *Salmonella* scheme with Grape-Tree for genomes within A) HC20_1348 and B) HC20_1487. 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=45423 and https://enterobase.warwick.ac.uk/ms_tree?tree_id=45424.