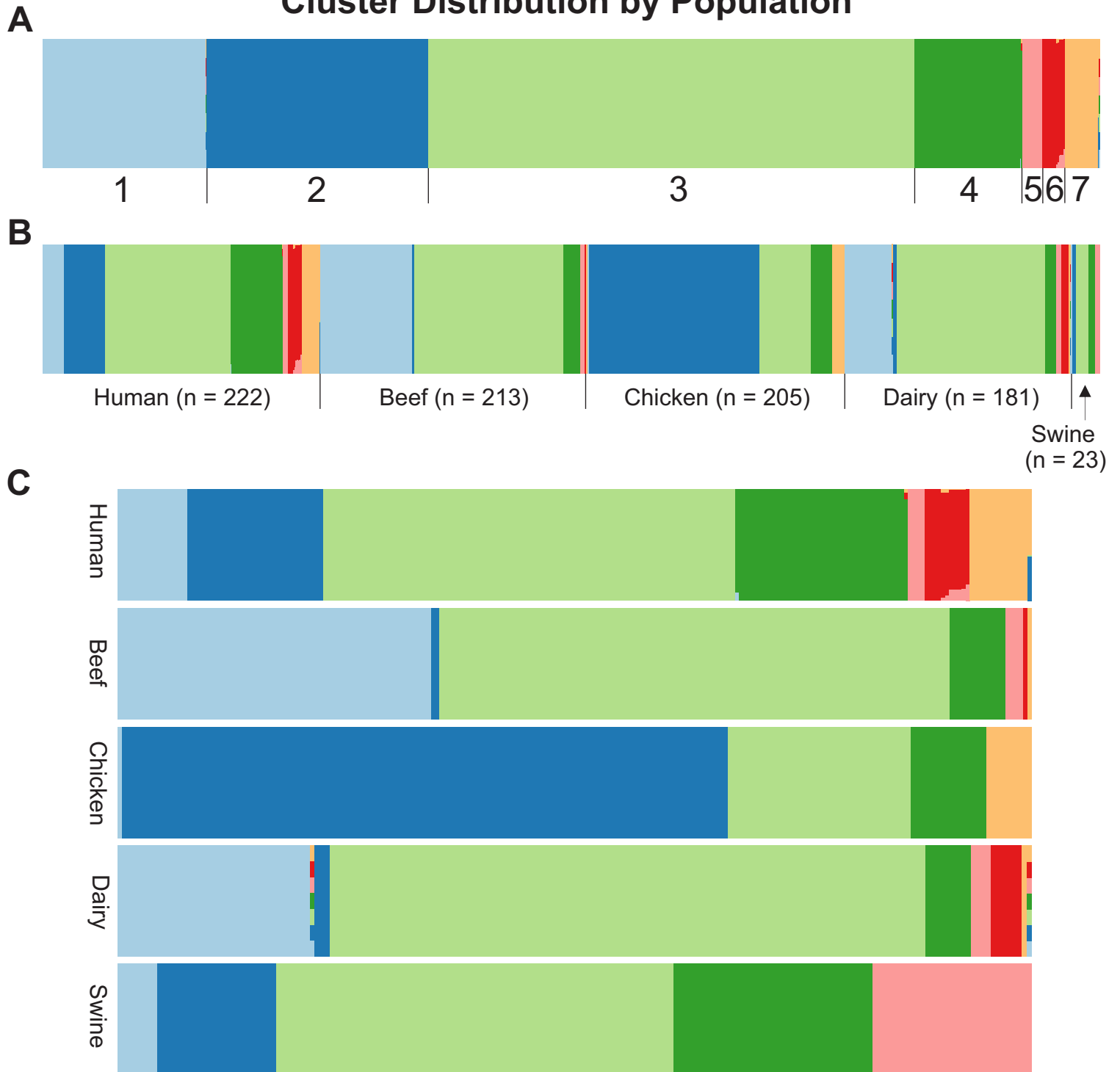Figure S1: Distribution of conserved population clusters as determined by STRUCTURE analysis of cgMLST allele profiles for 844 *C. jejuni* isolates. Cluster frequencies are shown both by A) aggregate population and B) source animal contribution. C) Proportional contribution of clusters illustrates differences in population composition between animal sources.