

Figure S1. Maximum likelihood tree of 80 *V. vulnificus* isolates. BS are based on 1000 replicates. The nine isolates of the present study are marked with orange.

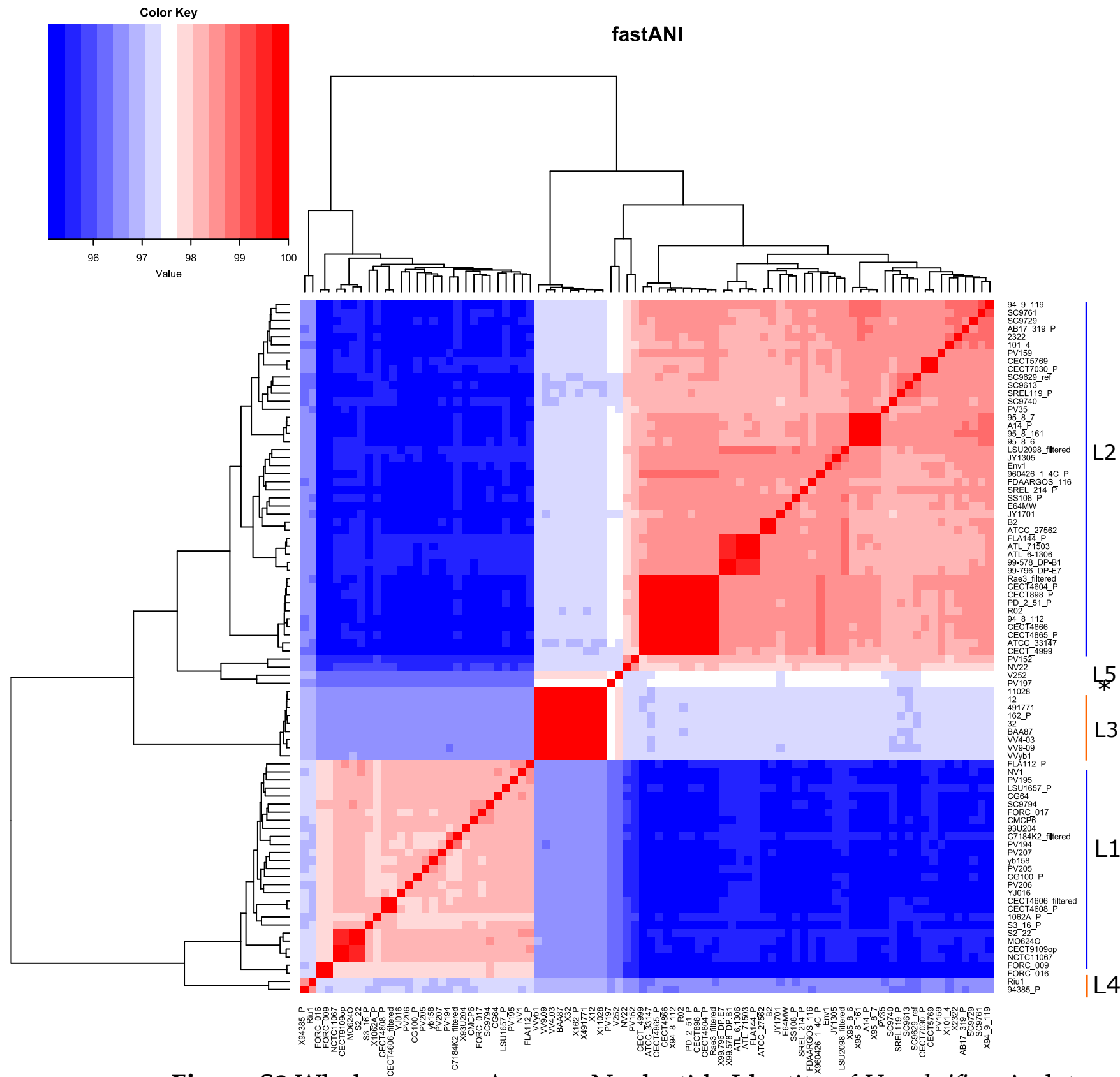


Figure S2. Whole-genome Average Nucleotide Identity of *V. vulnificus* isolates