

Supplementary Information of “Comparative Genomic Analysis Reveals the Potential Risk of *Vibrio parahaemolyticus* Isolated from Ready-to-eat Foods in China”

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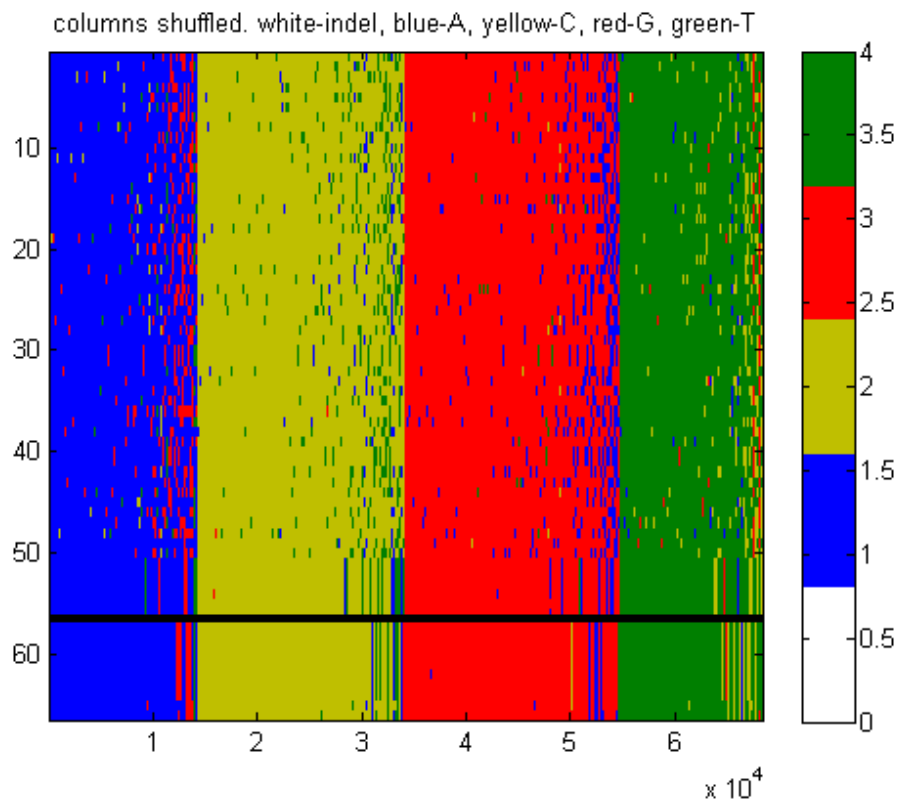


Figure S1. Bayesian Analysis of Population Structure of 66 global isolates of *Vibrio parahaemolyticus*. Each row in the figure is a strain and each column is a SNP, and horizontal black lines are drawn to separate the two clusters. The colour pattern of these SNPs within the black lines is consistent with them being assigned to the same BAPS group.

BIOFILM FORMATION - VIBRIO CHOLERAE

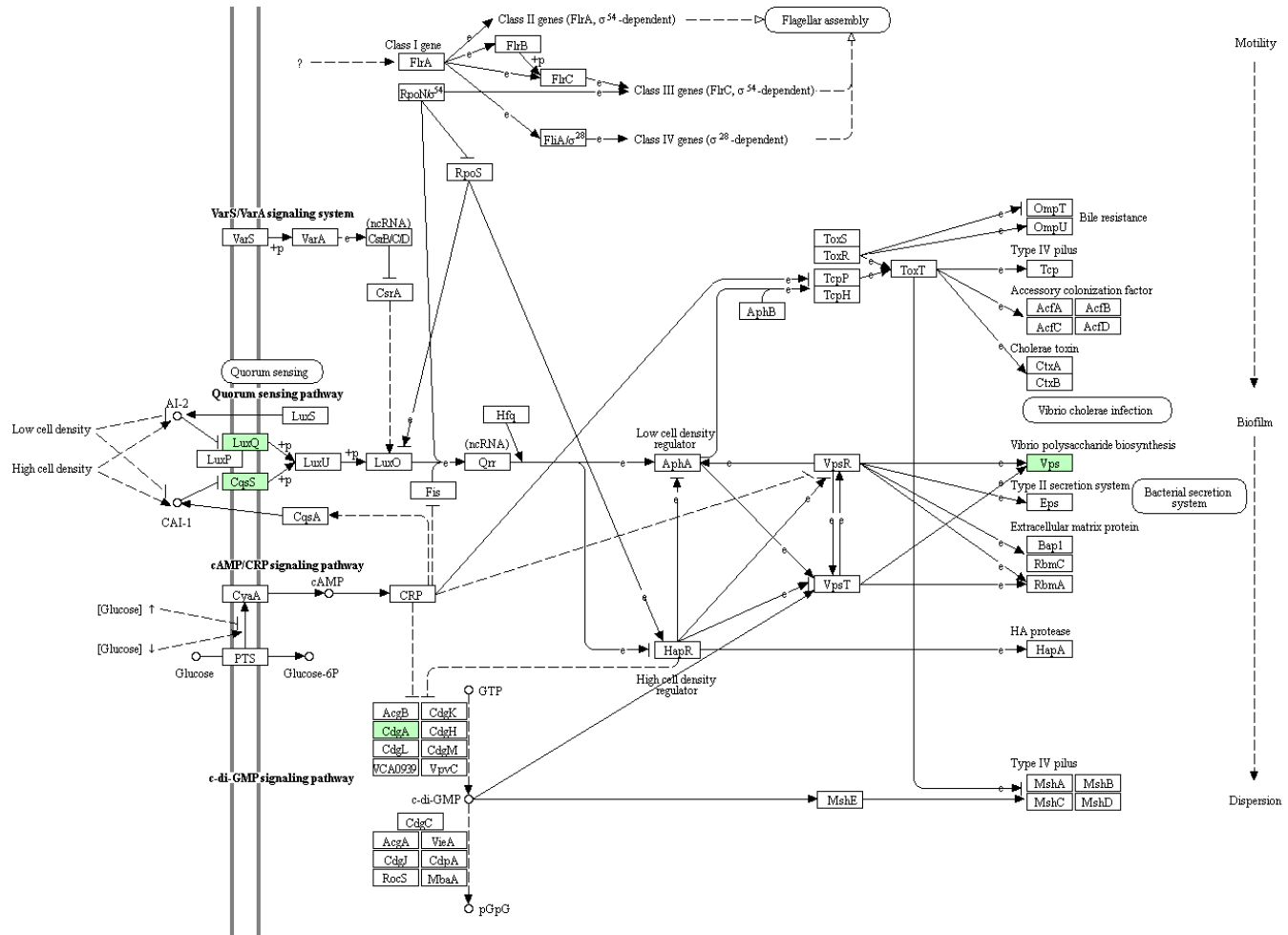


Figure S2. Pathway of biofilm formation (Ko05111). Green indicates the RTE food-related genes. LuxQ: group_2245 (K10909); cqsS: VPA0710 (K10916); exoP: group_261 (K16554); vpsM: group_6406 (K20920); cdgA: group_3579 (K20956).