S6 Fig. (A) Ternary comparison of flagellar gene distribution of six Arcobacter species with deep-branching species (Lebetimonas sp. JS138) and a close relative (Sulfurimonas gotlandica GD1). Below each ternary comparison, the flagellar genes of the Arcobacter species are depicted by ochre strips on linearized genomes, and the numerical symbols demonstrate two big flagellar gene clusters: Fcluster-1 and Fcluster-2. (B) The gene order of two big flagellar gene clusters (orange arrow) in Arcobacter species. The blue arrows represent two chemosensory genes cheYZ located inside of Fcluster-1.