



Figure S2:

Circular visualization map of all pangene categories and gene expression patterns (\log_2 ratio RPKM CDS:RPKM median) of (a) *A. salmonicida* LFI1238 and *V. natriegens* ATCC 14048 grown under (b) fast- and (c) slow-growing conditions. The areas marked in red correspond to “i” and “ii” in Fig. 2. Megabases are labeled as ticks. The genome comparison tool Circos [24] was used to create the figure.