



FIGURE S1. qRT-PCR to verify the RNA-seq data of strains A) HH01 / HH01 Δ jqsA and B) HH102 / HH102 Δ jqsA at early stationary phase. Shown are the normalized expressions of three technical and three (HH01/HH01 Δ jqsA) or two (HH102/HH102 Δ jqsA) biological samples. The graphs are shown with data relative to control using *rpoD* (HH01, HH102), *dnaG* (HH01) and *dnaB* (HH102, Table S2) as house keeping genes.