



Figure S2. RT-qPCR analysis by $\Delta\Delta\text{CT}$ method. Values normalized with *DhRPS3* gene and represented in \log_2 as fold changes of tRNA^{CAG}, aaRS, Trna^{Ser} and tRNA^{Leu} genes under different stress conditions. A) Expression profile of genes compared to each other under stress conditions tested in the logarithmic growth phase. B) Expression profile of genes compared to each other under stress conditions tested in the stationary growth phase. *p*-value: 0.01 (**), 0.001 (***), 0.0001 (****).