



Figure S2: Validation of the transcriptome data by q-RT-PCR analyses. The diagrams are representing the log₂-fold changes of the gene expression of SO_2500, SO_2660 and SO_3376 in surface-associated cells under hydrodynamic conditions (harvested after 0.25, 1, 2 and 4 hours) compared to planktonic culture. Plotted on the x-axis are the data values of the microarray analyses, on the y-axis the data values of the q-RT-PCR are shown. The coefficient of determination R^2 represents the statistical variance of the transcriptome data.