

Table S2. RT-PCR analysis of usher genes expression in the $\Delta mrkC$ mutant. Results are expressed in fold change expression (\pm SD) in the $\Delta mrkC$ mutant compared to the wild-type strain. Expression levels were compared by nonparametric one-way ANOVA; $p > 0.05$ (ANOVA)

target genes	fold expression (\pm SD) compared to the wild-type strain
<i>kpaC</i>	1.6 \pm 0.4
<i>kpbC</i>	2.3 \pm 0.5
<i>kpdC</i>	2.1 \pm 0.7
<i>kpeC</i>	1.8 \pm 0.7
<i>kpgC</i>	1.3 \pm 0.4
<i>kpjC</i>	1.9 \pm 0.6
<i>fimC</i>	1.5 \pm 0.7