

Table S2 - Levels of *mepA* expression measured by quantitative real-time PCR. *In vitro*-selected strains were normalized to SA43. Mean expression values and standard deviations were calculated from triplicates of two independent experiments. Statically significant variations in expression were calculated with ANOVA ($P < 0.05$).

Strain	Mean expression	Standard deviation	Significant $P < 0.05$
SA43	1.00	0.00	-
A2	4.11	0.84	NS
A5	199.82	13.40	***
A7	236.18	31.51	***
A10	178.50	19.73	***
B2	2.01	0.24	NS
B6	79.08	4.94	*
B7	298.32	38.08	***
B10	190.82	18.33	***
C2	2.10	0.30	NS
C4	211.92	14.76	***
C6	263.37	42.74	***
C7	45.22	0.72	NS