

Table S1. ANI values among tested strains

	<i>S. aureus</i> 35 G	<i>S. aureus</i> 36 G	<i>S. aureus</i> 39 G	<i>S. aureus</i> 45 G	<i>S. aureus</i> 47G
<i>S. aureus</i> 35 G	100	98.84	99.96	98.83	98.85
<i>S. aureus</i> 36 G	98.84	100	98.85	98.91	98.78
<i>S. aureus</i> 39 G	99.96	98.85	100	98.87	98.96
<i>S. aureus</i> 45 G	98.83	98.91	98.87	100	98.88
<i>S. aureus</i> 47 G	98.85	98.78	98.96	98.88	100

Table S2. Statistical significance between expressions after exposure to tested stressor among each strains based on Kruskal-Wallis ANOVA test.

stress factors	35G	39G	36G	45G	47G
pH=9.6	0.00273	0.001872	na	na	0.00511
pH=4.5	0.002808	0.001872	na	na	0.003129
4.5% NaCl	0.002063	0.002041	na	na	0.00267

na- not applicable , $p<0.05$ **Table S3.** Statistical significance between expressions each genes after exposure to tested stressor based on Kruskal-Wallis ANOVA test.

Gene	<i>p</i> -value
<i>sea</i>	0.06081
<i>seg</i>	0.019734*
<i>sei</i>	0.003207*
<i>selm</i>	0.000288*
<i>seln</i>	0.024283*
<i>selo</i>	0.000215*
<i>selu</i>	0.101489
<i>selp</i>	0.84896

*significant differences, $p<0.05$