

Table S1. Genes commonly modulated by the translocase blockage

Gene*	Annotated function	Regulation	Strain
<i>Nitrogen/amino acid metabolism</i>			
SLI0218	<i>narJ2</i> , nitrate reductase delta chain	Down	<i>sipY</i> and <i>secG</i> mutant
SLI0219	<i>narI2</i> , nitrate reductase gamma chain	Down	<i>sipY</i> and <i>secG</i> mutant
SLI1454	Possible amino oxidase	Up	<i>sipY</i> and <i>secG</i> mutant
SLI1498	<i>aroE</i> , shikimate 5-dehydrogenase	Down	<i>sipY</i> and <i>secG</i> mutant
SLI3073	<i>hutU</i> , urocanate hydratase	Up	<i>secG</i> and <i>lsp</i> mutant
<i>Oxidative stress/osmotic stress</i>			
SLI1866	Putative condensing enzyme	Up	<i>sipY</i> and <i>lsp</i> mutant
SLI5254	<i>sodN</i> , superoxide dismutase	Down	<i>sipY</i> and <i>lsp</i> mutant
<i>Membrane/transport protein</i>			
SLI5123	Putative small membrane protein	Down	<i>sipY</i> and <i>lsp</i> mutant
<i>Morphological differentiation</i>			
SLI0381	Possible glycosyl transferase	Down	<i>sipY</i> and <i>secG</i> mutant
SLI0382	Probable UDP-glucose/GDP-mannose family dehydrogenase (putative secreted protein)	Down	<i>sipY</i> and <i>secG</i> mutant
SLI0383	Hypothetical protein (possible target for <i>bldA</i> regulation)	Down	<i>sipY</i> and <i>secG</i> mutant
SLI0392	Possible methyltransferase similar to <i>S. peucetis</i> daunorubicin/ doxorubicin enzyme DnrX	Down	<i>sipY</i> and <i>secG</i> mutant
SLI0393	Possible transferase	Down	<i>sipY</i> and <i>secG</i> mutant
SLI0712	<i>lipR</i> putative transcriptional activator	Down	<i>sipY</i> and <i>secG</i> mutant
SLI1630	<i>cvnA9</i> , <i>rarA</i> putative integral membrane protein	Down	<i>sipY</i> and <i>secG</i> mutant
SLI1674	<i>chpC</i> , possible small membrane protein	Down	<i>sipY</i> and <i>secG</i> mutant
SLI1675	<i>chpH</i> , possible small membrane protein	Down	<i>sipY</i> and <i>secG</i> mutant
SLI1800	<i>chpE</i> , possible small membrane protein	Down	<i>sipY</i> and <i>secG</i> mutant
SLI2699	<i>chpG</i> , putative small membrane protein	Down	<i>sipY</i> and <i>secG</i> mutant
SLI2705	<i>chpF</i> , possible membrane protein	Down	<i>sipY</i> and <i>secG</i> mutant
SLI2717	<i>chpD</i> , putative small membrane protein	Down	<i>sipY</i> and <i>secG</i> mutant
SLI3323	<i>bldN</i> , probable RNA polymerase sigma factor	Down	<i>sipY</i> and <i>secG</i> mutant
SLI3549	<i>bldG</i> , probable anti-sigma factor antagonist	Down	<i>sipY</i> and <i>secG</i> mutant
SLI4768	<i>bldM</i> , probable two-component regulator	Down	<i>sipY</i> and <i>secG</i> mutant
SLI5113	<i>bldKB</i> , putative ABC transport system lipoprotein	Up	<i>sipY</i> and <i>secG</i> mutant
SLI5114	<i>bldKC</i> , putative ABC transport system integral membrane protein	Up	<i>sipY</i> and <i>secG</i> mutant
<i>Glycerol metabolism and glycolysis</i>			
SLI1659	<i>glpF</i> , probable glycerol uptake facilitator protein	Up	<i>sipY</i> and <i>secG</i> mutant
SLI1660	<i>glpK</i> , glycerol kinase	Up	<i>sipY</i> and <i>secG</i> mutant
<i>Secondary metabolites</i>			
SLI6073	<i>geoA</i> , possible cyclase germacradienol/ geosmin synthase	Down	<i>sipY</i> and <i>secG</i> mutant
<i>Secreted proteins</i>			
SLI0297	Possible secreted protein	Down	<i>sipY</i> , <i>secG</i> , <i>lsp</i> mutant
SLI0736	Possible secreted protein	Down	<i>secG</i> and <i>lsp</i> mutant
SLI0762	<i>sti1</i> , protease inhibitor	Down	<i>sipY</i> , <i>secG</i> , <i>lsp</i> mutant
SLI1860	Possible secreted protein	Down	<i>sipY</i> , <i>secG</i> , <i>lsp</i> mutant

SLI2116	Possible secreted protein	Down	<i>secG</i> and <i>lsp</i> mutant
SLI6197	Possible secreted protein	Down	<i>sipY</i> and <i>lsp</i> mutant
SLI6198	Possible secreted protein	Down	<i>sipY</i> and <i>lsp</i> mutant
SLI7657	Possible secreted protein	Down	<i>sipY</i> , <i>secG</i> , <i>lsp</i> mutant
<i>Other genes</i>			
SLI0682	Hypothetical protein	Down	<i>sipY</i> and <i>secG</i> mutant
SLI2822	cvnD12, posible conserverd ATP/GTP-binding protein	Down	<i>sipY</i> and <i>lsp</i> mutant
SLI6393	Probable transposase IS21/IS1162 family	Down	<i>sipY</i> and <i>secG</i> mutant
SLI7658	Hypothetical protein	Down	<i>sipY</i> and <i>lsp</i> mutant

*Genes names are as appeared in references [64,73].