

Table S1: Blast results of alignment of 16s rRNA nucleotide sequences

Bacterial stains	Description	Scientific Name	Accession
Susceptible. 6	<i>Staphylococcus saprophyticus</i> strain SUSCEPTIBLE-06 16S ribosomal RNA gene, partial sequence	<i>Staphylococcus saprophyticus</i>	OR050578.1
Susceptible. 8	<i>Staphylococcus pasteurii</i> strain SUSCEPTIBLE-08 16S ribosomal RNA gene, partial sequence	<i>Staphylococcus pasteurii</i>	OR050579.1
Susceptible. 16	<i>Enterobacter hormaechei</i> strain SUSCEPTIBLE-16 16S ribosomal RNA gene, partial sequence	<i>Enterobacter hormaechei</i>	OR050580.1
Susceptible. 46	<i>Bacillus toyonensis</i> strain SUSCEPTIBLE-46 16S ribosomal RNA gene, partial sequence	<i>Bacillus toyonensis</i>	OR050581.1
Tolerant. 30	<i>Bacillus thuringiensis</i> strain TOLERANT-30 16S ribosomal RNA gene, partial sequence	<i>Bacillus thuringiensis</i>	OR002134.1
Tolerant. 42	<i>Enterococcus gallinarum</i> strain TOLERANT-42 16S ribosomal RNA gene, partial sequence	<i>Enterococcus gallinarum</i>	OR002135.1
Tolerant. 52	<i>Staphylococcus saprophyticus</i> strain TOLERANT-52 16S ribosomal RNA gene, partial sequence	<i>Staphylococcus saprophyticus</i>	OR002133.1
Tolerant. 72	<i>Bacillus safensis</i> strain TOLERANT-72 16S ribosomal RNA gene, partial sequence	<i>Bacillus safensis</i>	OR002132.1
Tolerant. 104	<i>Enterococcus mundtii</i> strain TOLERANT-104 16S ribosomal RNA gene, partial sequence	<i>Enterococcus mundtii</i>	OR002136.1