

Additional file 3: Table S3. Identity of bacteria isolated from the Stimpson Wild Life Sanctuary's soils with various salt levels.
Bacterial strains were identified based on the 16S rRNA sequence.

Sample Name	GenBank ID	Description	Query cover	e value
<u>0 PPT</u>				
8G/3E	gb KU937390.1	<i>Enterococcus faecalis</i> strain GX27	99%	0
8B	gb GQ337884.1	<i>Enterococcus faecalis</i> strain KLDS4.0341	99%	0
*6D/12G	gb EU794735.1	<i>Enterococcus</i> sp. Le5-1a	100%	0
*3F	gb KC213478.1	<i>Enterococcus durans</i> strain GM18	100%	0
<u>14ppt</u>				
8F/1H	gb KU937390.1	<i>Enterococcus faecalis</i> strain GX27	99%	0
1G/8D	gb GQ337884.1	<i>Enterococcus faecalis</i> strain KLDS4.0341	99%	0
7C	gb KU937390.1	<i>Enterococcus faecalis</i> strain GX27	99%	0
7A	gb KT343158.1	<i>Enterococcus faecalis</i> strain JF85	100%	0
3B	gb EU708623.1	<i>Enterococcus faecalis</i> strain XR7	99%	0
9D	gb KP241784.1	<i>Serratia</i> sp. SH-AB-1	99%	0
*2E	gb HQ831381.1	<i>Enterococcus faecalis</i> strain Na15	99%	0
*2F	gb HQ184922.1	<i>Enterococcus faecalis</i> strain FUA3334	99%	0
*7B	gb JX006538.1	Bacterium NLAE-zl-H323	99%	0
<u>20ppt</u>				
*6B/8h/10F	gb KT343158.1	<i>Enterococcus faecalis</i> strain JF85	100%	0
<u>50ppt</u>				
5G/5H	gb EU708623.1	<i>Enterococcus faecalis</i> strain XR7	99%	0
9C/12C	gb KP241784.1	<i>Serratia</i> sp. SH-AB-1	99%	0
5E	gb KX752874.1	<i>Enterococcus hirae</i> strain SNNU0253	99%	0
*5D	gb JX006721.1	Bacterium NLAE-zl-H515	99%	0
*3G	gb HQ831431.1	<i>Enterococcus faecalis</i> strain TLME3	99%	0
*5F	gb JF947360.1	<i>Morganella morganii</i> strain 2113	98%	0
*9B	gb KT894728.1	<i>Serratia marcescens</i> strain JND-KHCo-24B	99%	0
*10H	gb AF513469.1	Enterobacteriaceae bacterium PH31	99%	0