

Supplementary figures

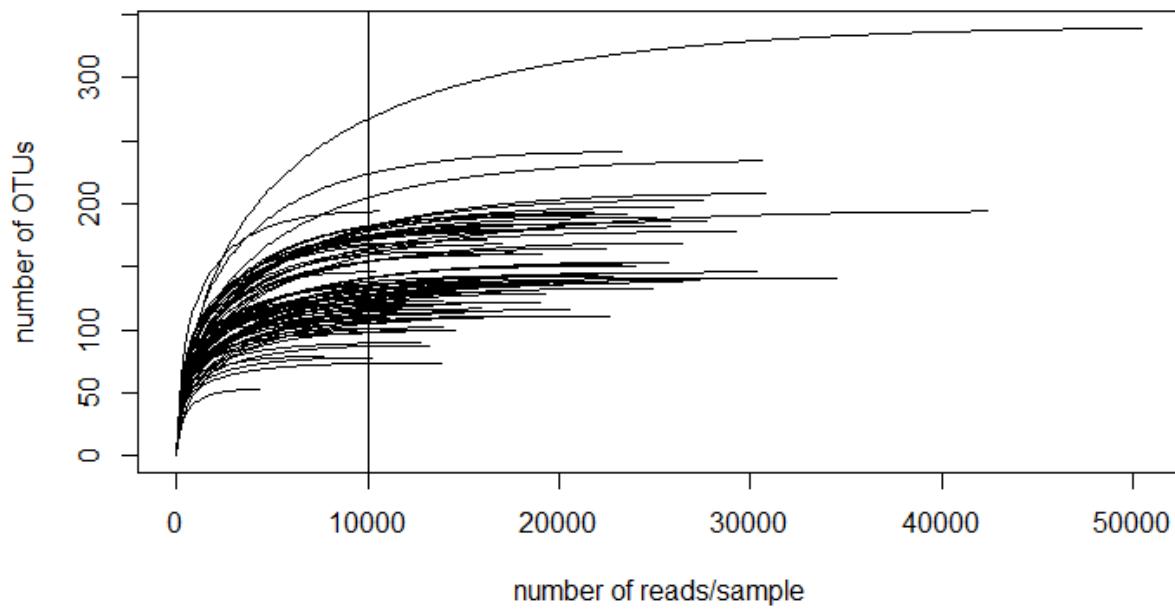
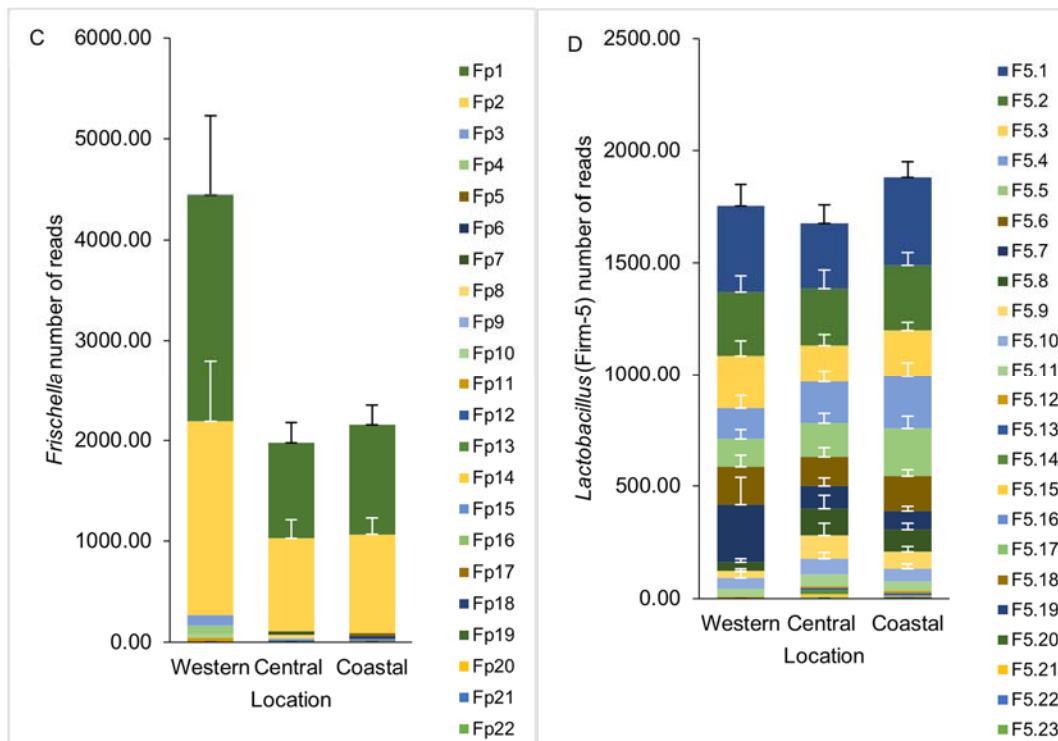
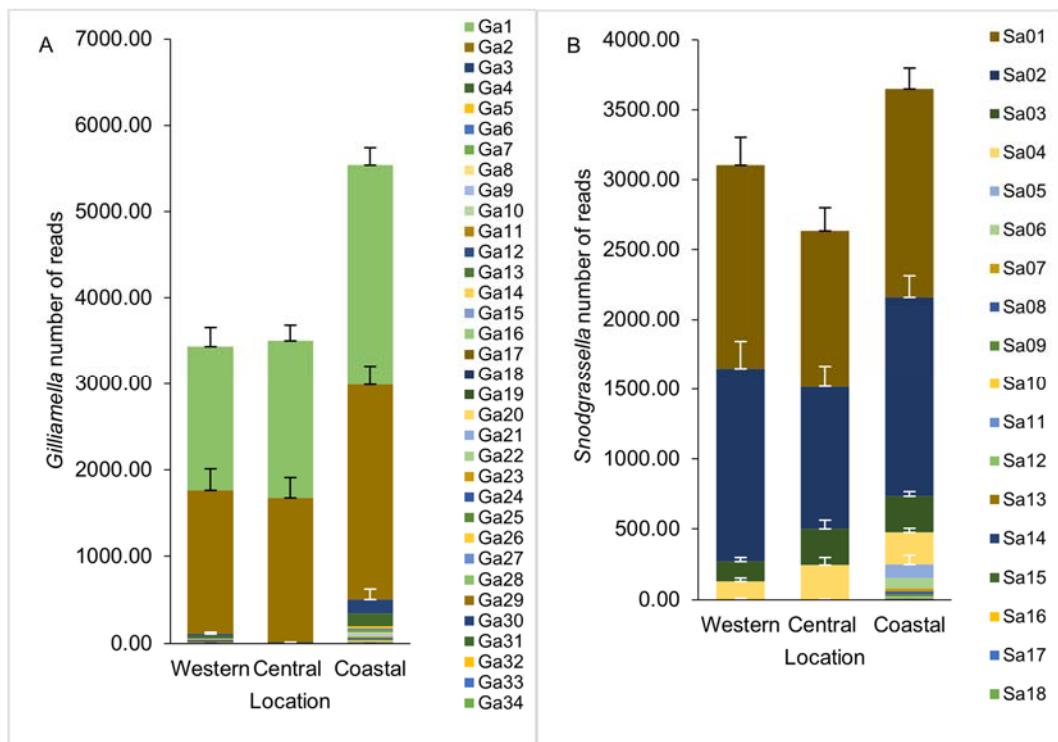
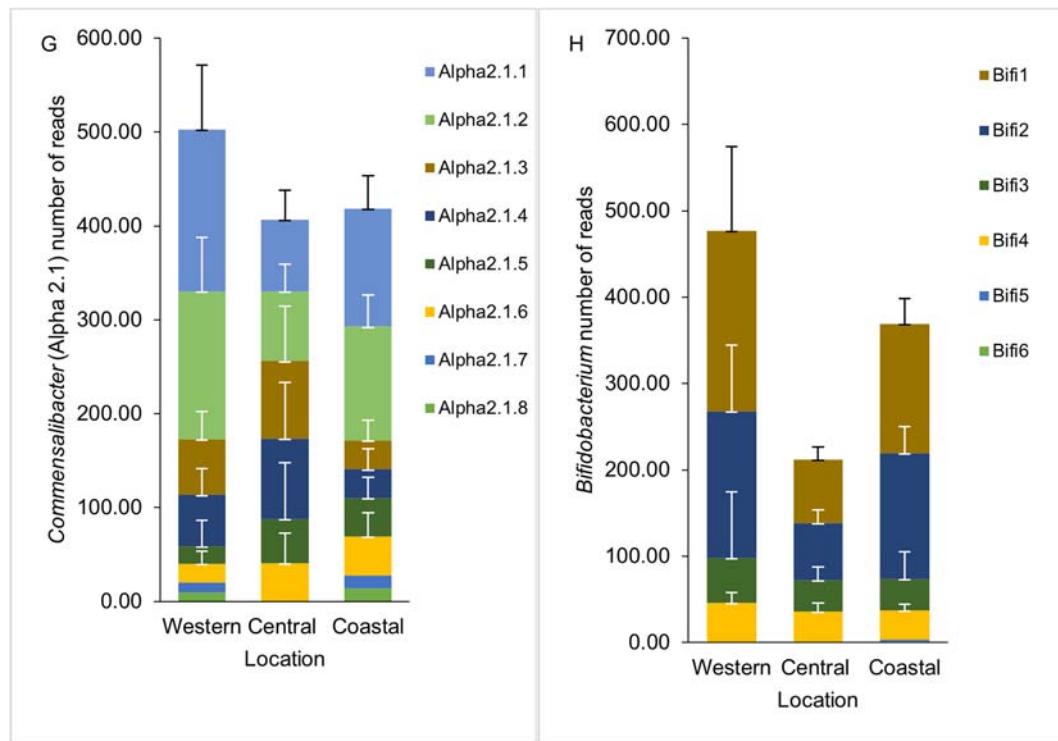
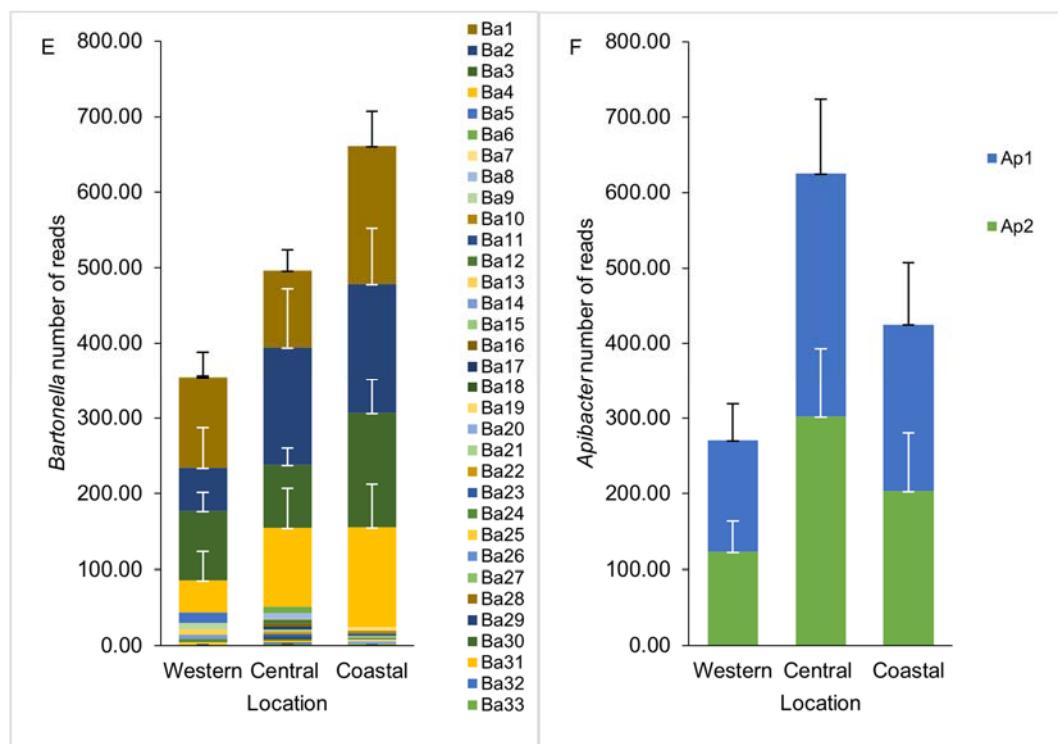


Figure S1: Rarefaction curve (OTUs per reads/samples). The rarefaction value is 10,000 reads.





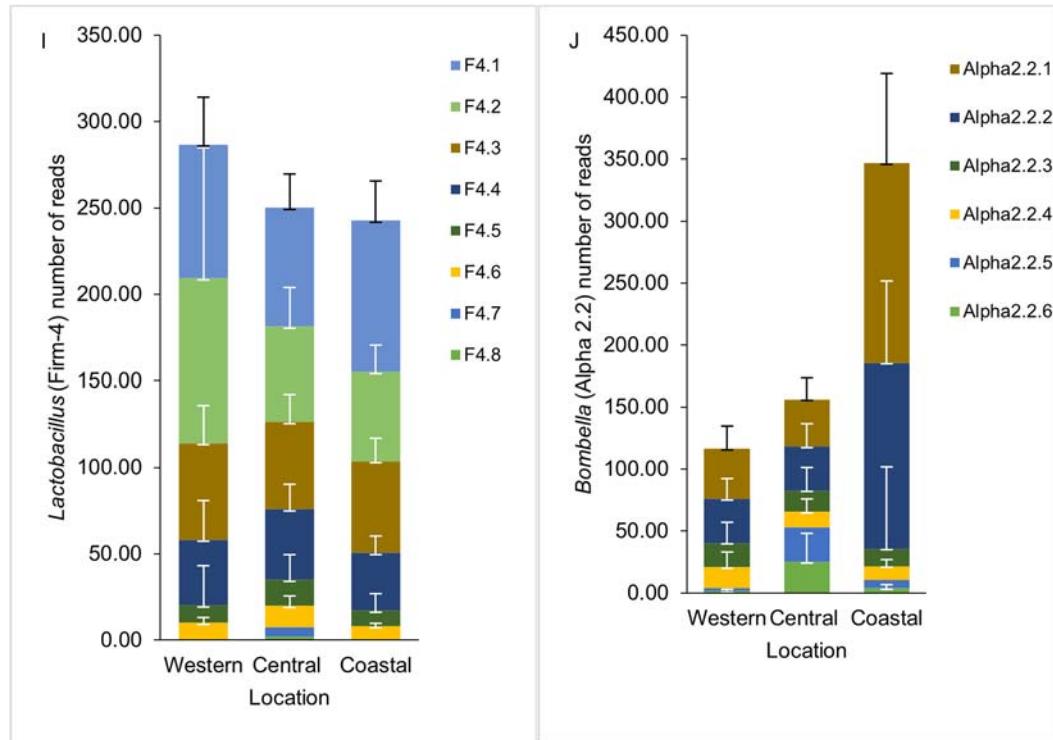


Figure S2: Amplicon sequence variant (ASV) relative abundance of the main bacteria core members.

Table S2: Most abundant environmental bacterial abundance and prevalence.

| Environmental bacterial genus | Environmental bacterial abundance (%) and prevalence (%) | | | | | |
|-------------------------------|--|------------|-----------|------------|-----------|------------|
| | Western | | Central | | Coastal | |
| | Abundance | Prevalence | Abundance | Prevalence | Abundance | Prevalence |
| <i>Pseudomonas</i> | 3.1 | 87.2 | 1.2 | 77.9 | 2.3 | 62.8 |
| <i>Acidovorax</i> | 3.4 | 87.2 | 2.4 | 81.4 | 0.6 | 61.6 |
| <i>Spiroplasma</i> | 2.2 | 83.7 | 3.9 | 76.7 | 0.0 | 54.7 |
| <i>Gluconobacter</i> | 1.9 | 93.0 | 1.5 | 90.7 | 1.8 | 89.5 |
| <i>Acinetobacter</i> | 0.0 | 79.1 | 3.8 | 75.6 | 0.0 | 55.8 |
| <i>Klebsiella</i> | 1.2 | 86.0 | 0.6 | 79.1 | 0.7 | 64.0 |
| <i>Fructobacillus</i> | 1.4 | 88.4 | 0.8 | 84.9 | 0.3 | 70.9 |
| <i>Enterobacter</i> | 0.8 | 84.9 | 0.3 | 74.4 | 0.4 | 58.1 |
| <i>Weissella</i> | 1.8 | 82.6 | 0.1 | 73.3 | 0.0 | 52.3 |
| <i>Tatumella</i> | 0.6 | 90.7 | 0.5 | 87.2 | 0.3 | 77.9 |

Table S3: PERMANOVA analysis showed no bacterial community variation among locations. $R^2 = 0.03074$, $P = 0.198$.

| Group 1 | Group 2 | F.Model | R ² | p-value | p-adjusted |
|---------|---------|----------|----------------|---------|------------|
| Western | Coastal | 1.378317 | 0.02282801 | 0.136 | 0.408 |
| Western | Central | 1.110891 | 0.02638013 | 0.346 | 1.000 |
| Coastal | Central | 1.570429 | 0.02324137 | 0.060 | 0.180 |

Table S4: Pairwise comparisons of bacterial relative abundance showed significant variation of *Gilliamella* among locations. Statistical significance are indicated with different letters.

| Bacterial genus | Pairwise Comparison | | | Western | Central | Coastal |
|-------------------------------------|---------------------|---------|---|---------|---------|---------|
| | F-value | P-value | | | | |
| <i>Gilliamella</i> | 3.105 | 0.0501 | b | ab | a | |
| <i>Snodgrassella</i> | 0.18 | 0.836 | a | a | a | |
| <i>Frischella</i> | 2.156 | 0.122 | a | a | a | |
| <i>Lactobacillus</i> (Firm-5) | 0.675 | 0.512 | a | a | a | |
| <i>Bartonella</i> | 0.38 | 0.685 | a | a | a | |
| <i>Apibacter</i> | 1.756 | 0.179 | a | a | a | |
| <i>Commensalibacter</i> (Alpha 2.1) | 0.908 | 0.407 | a | a | a | |
| <i>Bifidobacter</i> | 1.821 | 0.168 | a | a | a | |
| <i>Lactobacillus</i> (Firm-4) | 0.776 | 0.464 | a | a | a | |
| <i>Bombella</i> (Alpha 2.2) | 0.461 | 0.632 | a | a | a | |