

Supplementary Table 5: Genera achieving a nominal significance level of $P<0.05$ in four-way comparison.

| Location | Phylum | Genus | FET P-value | FET P-value (corr) | KWT P-value | KWT P-value (corr) |
|---------------|----------------|-------------------------------------|----------------------|----------------------|----------------------|----------------------|
| Pouch | Actinobacteria | <i>Corynebacterium</i> | 0.02 | 0.15 | 0.03 | 0.16 |
| | Bacteroidetes | <i>Bacteroides</i> | 3.6×10^{-6} | 3.0×10^{-4} | 5.7×10^{-7} | 4.7×10^{-5} |
| | | <i>Parabacteroides</i> | 1.2×10^{-4} | 4.9×10^{-3} | 7.8×10^{-4} | 0.02 |
| | Firmicutes | <i>Moryella</i> | 2.5×10^{-3} | 0.05 | 1.4×10^{-3} | 0.03 |
| | | <i>Blautia</i> | 3.0×10^{-3} | 0.05 | 2.3×10^{-4} | 9.5×10^{-3} |
| | | <i>Dorea</i> | 0.02 | 0.13 | 0.01 | 0.09 |
| | | <i>Phascolarctobacterium</i> | 9.6×10^{-3} | 0.10 | 0.02 | 0.15 |
| | | IS <i>Erysipelotrichaceae</i> | 0.02 | 0.14 | 0.03 | 0.16 |
| | | <i>Dialister</i> | 0.02 | 0.13 | 0.02 | 0.15 |
| | | Uncultured <i>Lachnospiraceae</i> | 0.04 | 0.20 | 6.9×10^{-3} | 0.07 |
| | | Uncultured <i>Ruminococcaceae</i> | 0.04 | 0.20 | 7.1×10^{-3} | 0.07 |
| | | <i>Subdoligranulum</i> | 0.01 | 0.13 | 0.03 | 0.16 |
| | | Uncultured Family XIII IS | 8.5×10^{-3} | 0.10 | 7.7×10^{-3} | 0.07 |
| | | <i>Roseburia</i> | 0.07 | 0.31 | 0.04 | 0.17 |
| | Proteobacteria | <i>Sutterella</i> | 2.5×10^{-3} | 0.05 | 2.4×10^{-3} | 0.04 |
| | | <i>Rhodobacter</i> | 0.01 | 0.09 | 0.03 | 0.16 |
| | | <i>Haemophilus</i> | 0.13 | 0.39 | 5.9×10^{-3} | 0.12 |
| Afferent Limb | Actinobacteria | Uncultured <i>Coriobacteriaceae</i> | 0.03 | 0.17 | 0.03 | 0.20 |
| | | <i>Collinsella</i> | 0.02 | 0.16 | 0.02 | 0.19 |
| | Bacteroidetes | <i>Bacteroides</i> | 6.9×10^{-6} | 2.9×10^{-4} | 4.2×10^{-6} | 3.4×10^{-4} |
| | | <i>Parabacteroides</i> | 2.2×10^{-3} | 0.03 | 8.8×10^{-3} | 0.10 |
| | Firmicutes | <i>Moryella</i> | 0.03 | 0.17 | 0.03 | 0.20 |
| | | <i>Blautia</i> | 8.2×10^{-7} | 6.8×10^{-5} | 1.1×10^{-4} | 4.7×10^{-3} |
| | | <i>Dorea</i> | 1.0×10^{-3} | 0.03 | 4.2×10^{-3} | 0.07 |
| | | <i>Anaerococcus</i> | 2.8×10^{-3} | 0.05 | 8.4×10^{-3} | 0.11 |
| | | IS <i>Erysipelotrichaceae</i> | 4.5×10^{-3} | 0.05 | 4.0×10^{-3} | 0.07 |
| | | <i>Roseburia</i> | 0.02 | 0.15 | 0.08 | 0.30 |
| | | Uncultured Family XIII IS | 0.01 | 0.13 | 0.03 | 0.19 |
| | | <i>Granulicatella</i> | 0.07 | 0.31 | 0.03 | 0.19 |
| | | <i>Dialister</i> | 0.05 | 0.23 | 0.05 | 0.21 |
| | | <i>Lactobacillus</i> | 0.03 | 0.17 | 0.01 | 0.15 |
| | | <i>Mogibacterium</i> | 0.04 | 0.20 | 0.05 | 0.21 |
| | | <i>Phascolarctobacterium</i> | 0.02 | 0.16 | 0.03 | 0.19 |
| | Proteobacteria | <i>Sutterella</i> | 1.7×10^{-3} | 0.03 | 8.5×10^{-4} | 0.02 |
| | | <i>Rhodobacter</i> | 0.03 | 0.17 | 0.05 | 0.21 |
| | | <i>Haemophilus</i> | 0.12 | 0.37 | 0.05 | 0.21 |

FET=Fisher's exact test (dichotomous results); KWT=Kruskal-Wallis Test (continuous results). Highlighted rows are those which remained significant in either analysis after FDR correction for multiple testing.