

Table 2. Bacterial taxa that differ between IBS, IBS+bloat, IBS-bloat, and healthy controls

| Y Variable | Group | Mean Abundance* | Std Dev. | Std Err. | N | P ANOVA | P Tukey |
|--------------------------------------|--------------|-----------------|----------|----------|----|---------|---------|
| Lactobacillus | IBS Bloat | -3.215 | 0.599 | 0.084 | 51 | 0 | |
| Lactobacillus | IBS No Bloat | -2.995 | 1.465 | 0.598 | 6 | 0 | 0 |
| Lactobacillus | HC | -4.283 | 0.344 | 0.092 | 14 | 0 | 0.001 |
| Anaerotruncus | IBS Bloat | -3.479 | 0.563 | 0.079 | 51 | 0.007 | 0.01 |
| Anaerotruncus | IBS No Bloat | -4.198 | 0.369 | 0.151 | 6 | 0.007 | |
| Anaerotruncus | HC | -3.767 | 0.571 | 0.153 | 14 | 0.007 | |
| Streptococcus thermophilus LMG 18311 | IBS Bloat | -2.379 | 0.692 | 0.097 | 51 | 0.007 | |
| Streptococcus thermophilus LMG 18311 | IBS No Bloat | -1.887 | 0.567 | 0.232 | 6 | 0.007 | 0.034 |
| Streptococcus thermophilus LMG 18311 | HC | -2.913 | 0.746 | 0.199 | 14 | 0.007 | 0.01 |
| Ruminococcaceae | IBS Bloat | -0.954 | 0.584 | 0.082 | 51 | 0.009 | 0.011 |
| Ruminococcaceae | IBS No Bloat | -1.703 | 1.02 | 0.416 | 6 | 0.009 | |
| Ruminococcaceae | HC | -0.841 | 0.242 | 0.065 | 14 | 0.009 | 0.009 |
| Streptococcus parasanguinis (T) | IBS Bloat | -3.381 | 0.671 | 0.094 | 51 | 0.011 | |
| Streptococcus parasanguinis (T) | IBS No Bloat | -2.861 | 0.912 | 0.372 | 6 | 0.011 | 0.071 |
| Streptococcus parasanguinis (T) | HC | -3.843 | 0.623 | 0.166 | 14 | 0.011 | 0.012 |
| Streptococcus | IBS Bloat | -2.255 | 0.642 | 0.09 | 51 | 0.013 | |
| Streptococcus | IBS No Bloat | -1.776 | 0.546 | 0.223 | 6 | 0.013 | 0.078 |
| Streptococcus | HC | -2.679 | 0.653 | 0.175 | 14 | 0.013 | 0.014 |
| Subdoligranulum variabile DSM 15176 | IBS Bloat | -2.047 | 1.115 | 0.156 | 51 | 0.023 | 0.017 |
| Subdoligranulum variabile DSM | IBS No Bloat | -3.4 | 1.23 | 0.502 | 6 | 0.023 | |

| | | | | | | | |
|-------------------------------------|--------------|--------|-------|-------|----|-------|-------|
| Subdoligranulum variabile DSM 15176 | HC | -2.248 | 1.038 | 0.277 | 14 | 0.023 | |
| OTU4438 | IBS Bloat | -3.6 | 0.742 | 0.104 | 51 | 0.028 | |
| OTU4438 | IBS No Bloat | -4.059 | 0.659 | 0.269 | 6 | 0.028 | |
| OTU4438 | HC | -3.152 | 0.653 | 0.174 | 14 | 0.028 | 0.032 |
| OTU2915 | IBS Bloat | -3.199 | 0.916 | 0.128 | 51 | 0.03 | |
| OTU2915 | IBS No Bloat | -3.847 | 1.017 | 0.415 | 6 | 0.03 | |
| OTU2915 | HC | -2.682 | 0.838 | 0.224 | 14 | 0.03 | 0.028 |
| Anaerovorax_OTU14 | IBS Bloat | -3.432 | 0.608 | 0.085 | 51 | 0.036 | 0.028 |
| Anaerovorax_OTU14 | IBS No Bloat | -4.136 | 0.688 | 0.281 | 6 | 0.036 | |
| Anaerovorax_OTU14 | HC | -3.476 | 0.631 | 0.169 | 14 | 0.036 | |
| Oscillibacter_OTU4 | IBS Bloat | -2.704 | 0.73 | 0.102 | 51 | 0.052 | |
| Oscillibacter_OTU4 | IBS No Bloat | -3.369 | 0.909 | 0.371 | 6 | 0.052 | |
| Oscillibacter_OTU4 | HC | -2.494 | 0.626 | 0.167 | 14 | 0.052 | 0.042 |

--- IBS+B = IBS with Bloating; IBS-B = IBS without Bloating; HC=Healthy Controls

*mean of log10-transformed relative abundances