

S7 Table. RDP classification of the sequence reads at class level. Data are presented as the means \pm S.D and analyzed by Mann-Whitney test, *** $P<0.001$, ** $P<0.01$ and * $P<0.05$ vs the HFD group.

| Class | Control (%) | HFD (%) | HFD+M (%) | HFD+FO600 (%) | HFD+KO600 (%) | HFD+FO300KO300 (%) | HFD+FO400KO200 (%) | HFD+FO450KO150 (%) |
|-------------------------------|----------------------|------------------|----------------------|---------------------|----------------------|---------------------|----------------------|---------------------|
| <i>Bacteroidia</i> | 68.38 \pm 6.41↑*** | 33.77 \pm 4.88 | 43.16 \pm 8.74↑* | 53.9 \pm 10.37↑** | 43.73 \pm 6.89↑** | 34.53 \pm 6.88↑ | 39.78 \pm 8.89↑ | 32.75 \pm 5.67↓ |
| <i>Clostridia</i> | 16.46 \pm 3.16↓*** | 38.63 \pm 6.74 | 19.91 \pm 4.32↓*** | 35.14 \pm 7.62↓ | 21.29 \pm 3.41↓*** | 38.34 \pm 5.96↓ | 36.34 \pm 7.42↓ | 35.4 \pm 5.87↓ |
| <i>Delta proteobacteria</i> | 1.57 \pm 0.38↓*** | 11.95 \pm 2.65 | 5.67 \pm 1.32↓** | 1.35 \pm 0.23↓*** | 6.01 \pm 2.46↓** | 20.73 \pm 4.67↑** | 3.26 \pm 2.34↓*** | 9.74 \pm 4.67↓ |
| <i>Bacilli</i> | 1.17 \pm 0.24↓** | 6.42 \pm 1.17 | 8.63 \pm 2.13↑ | 2.38 \pm 0.34↓*** | 10.57 \pm 3.11↑* | 0.69 \pm 0.12↓*** | 6.23 \pm 1.47↓ | 4.57 \pm 1.37↓ |
| <i>Erysipelotrichia</i> | 6.45 \pm 1.13↑** | 1.81 \pm 0.44 | 11.13 \pm 2.24↑*** | 1.01 \pm 0.18↓ | 4.44 \pm 1.56↑** | 1.71 \pm 0.22↓ | 4.65 \pm 1.33↑** | 4.56 \pm 1.42↑** |
| <i>Epsilon proteobacteria</i> | 1.69 \pm 0.32↓* | 2.59 \pm 0.32 | 2.8 \pm 0.43↑ | 2.21 \pm 0.28↓ | 2.15 \pm 0.98↓ | 1.19 \pm 0.07↓* | 0.87 \pm 0.16↓*** | 7.06 \pm 2.17↑*** |
| <i>Actinobacteria</i> | 2.17 \pm 0.47↑* | 1.74 \pm 0.27 | 4.29 \pm 0.26↑ | 1.25 \pm 0.16↓ | 5.1 \pm 1.17↑** | 0.79 \pm 0.04↓* | 3.65 \pm 0.78↑*** | 2.32 \pm 0.43↑** |
| <i>Gammaproteobacteria</i> | 0.52 \pm 0.11↑ | 0.47 \pm 0.13 | 2.06 \pm 0.17↑** | 0.48 \pm 0.24↑ | 1.36 \pm 0.05↑*** | 0.56 \pm 0.26↑*** | 0.96 \pm 0.22↑*** | 0.9 \pm 0.26↑*** |
| <i>Betaproteobacteria</i> | 0.8 \pm 0.16↑ | 0.59 \pm 0.24 | 0.77 \pm 0.06↑ | 0.43 \pm 0.17↓ | 1.5 \pm 0.17↑** | 0.21 \pm 0.01↓*** | 1.2 \pm 0.32↑*** | 0.7 \pm 0.12↑** |
| <i>Alphaproteobacteria</i> | 0.18 \pm 0.03↓ | 0.28 \pm 0.04 | 0.25 \pm 0.05↓ | 0.21 \pm 0.09↓ | 0.84 \pm 0.11↑* | 0.13 \pm 0.04↓* | 0.59 \pm 0.18↑*** | 0.36 \pm 0.11↑** |
| <i>Chloroplast</i> | 0.11 \pm 0.04↓* | 0.21 \pm 0.05 | 0.26 \pm 0.11↑ | 0.16 \pm 0.05↓ | 0.79 \pm 0.23↑** | 0.07 \pm 0.01↓** | 0.73 \pm 0.25↑*** | 0.36 \pm 0.14↑ |
| <i>Unclassified</i> | 0.15 \pm 0.01↓* | 0.59 \pm 0.03 | 0.25 \pm 0.03↓ | 0.5 \pm 0.07↓ | 0.17 \pm 0.06↓* | 0.38 \pm 0.02↓ | 0.25 \pm 0.11↓ | 0.2 \pm 0.07↓** |
| <i>Planctomycetia</i> | 0.09 \pm 0.01↓* | 0.18 \pm 0.07 | 0.29 \pm 0.09↑ | 0.14 \pm 0.03↓ | 0.59 \pm 0.08↑** | 0.06 \pm 0.01↓* | 0.47 \pm 0.14↑*** | 0.2 \pm 0.09↑** |
| <i>Deferribacteres</i> | 0.01 \pm 0.002↓** | 0.28 \pm 0.03 | 0.05 \pm 0.001↓* | 0.61 \pm 0.17↑** | 0.18 \pm 0.04↓ | 0.45 \pm 0.02↑*** | 0.02 \pm 0.005↓*** | 0.41 \pm 0.03↑*** |
| <i>Cyanobacteria</i> | 0.04 \pm 0.013↓ | 0.09 \pm 0.01 | 0.1 \pm 0.01↑ | 0.05 \pm 0.01↓ | 0.31 \pm 0.13↑** | 0.03 \pm 0.006↓ | 0.25 \pm 0.02↑ | 0.11 \pm 0.04↑ |
| <i>Spartobacteria</i> | 0.03 \pm 0.005↓ | 0.07 \pm 0.01 | 0.07 \pm 0.01 | 0.04 \pm 0.003↓ | 0.21 \pm 0.09↑ | 0.02 \pm 0.003↓ | 0.13 \pm 0.04↑ | 0.07 \pm 0.01↓ |
| <i>Sphingobacteriia</i> | 0.02 \pm 0.002↓ | 0.04 \pm 0.001 | 0.09 \pm 0.02↑ | 0.03 \pm 0.005↓ | 0.18 \pm 0.04↑* | 0.02 \pm 0.004↓* | 0.15 \pm 0.03↑*** | 0.07 \pm 0.02↑** |
| <i>Flavobacteriia</i> | 0.03 \pm 0.004↓ | 0.06 \pm 0.002 | 0.03 \pm 0.001↓ | 0.03 \pm 0.001↓ | 0.13 \pm 0.01↑** | 0.03 \pm 0.004↓* | 0.09 \pm 0.01↑* | 0.03 \pm 0.006* |
| <i>Phycisphaerae</i> | 0.03 \pm 0.005↓ | 0.04 \pm 0.001 | 0.04 \pm 0.002 | 0.02 \pm 0.001↓ | 0.09 \pm 0.02↑ | 0.02 \pm 0.002↓ | 0.08 \pm 0.02↑ | 0.02 \pm 0.002↓ |
| <i>Opitutae</i> | 0.02 \pm 0.003↓ | 0.03 \pm 0.003 | 0.02 \pm 0.001↓ | 0.01 \pm 0.001↓ | 0.07 \pm 0.01↑ | 0.04 \pm 0.006↓ | 0.09 \pm 0.01↑ | 0.03 \pm 0.004↑ |