

**Table S6.** RDP classification of the sequence ratio at phylum level. Data are represented as the means  $\pm$  S.D and analysed by Mann-Whitney test. \*\*\* $P<0.001$ , \*\* $P<0.01$  and \* $P<0.05$  vs the HFD group.

Phylum	Control (%)	HFD (%)	HFD+M (%)	HFD+FO600 (%)	HFD+KO600 (%)	HFD+FO300KO300 (%)	HFD+FO400KO200 (%)	HFD+FO450KO150 (%)
<i>Bacteroidetes</i>	68.44 $\pm$ 4.67↑***	33.88 $\pm$ 3.21	43.3 $\pm$ 4.22↑***	34.58 $\pm$ 4.21↑	53.96 $\pm$ 4.31↑***	44.11 $\pm$ 5.42↑***	40.06 $\pm$ 5.17↑**	32.87 $\pm$ 4.88↓
<i>Firmicutes</i>	24.13 $\pm$ 2.42↓***	46.93 $\pm$ 3.34	39.68 $\pm$ 3.17↓**	40.87 $\pm$ 3.47↓**	38.55 $\pm$ 8.75↓**	36.32 $\pm$ 3.41↓**	47.23 $\pm$ 3.43↑	44.58 $\pm$ 5.87↓
<i>Proteobacteria</i>	4.76 $\pm$ 1.22↓**	15.89 $\pm$ 2.33	11.56 $\pm$ 2.19↓*	22.83 $\pm$ 2.21↑**	4.67 $\pm$ 1.15↓**	11.87 $\pm$ 2.45↓**	6.9 $\pm$ 2.13↓**	18.76 $\pm$ 2.16↑*
<i>Actinobacteria</i>	2.17 $\pm$ 0.89↑	1.74 $\pm$ 0.67	4.29 $\pm$ 1.78↑*	0.79 $\pm$ 0.11↓**	1.25 $\pm$ 0.32↓	5.1 $\pm$ 1.88↑*	3.65 $\pm$ ↑*	2.32 $\pm$ 0.88↑
<i>Cyanobacteria/Chloroplast</i>	0.15 $\pm$ 0.04↓	0.3 $\pm$ 0.02	0.36 $\pm$ 0.13↑	0.1 $\pm$ 0.012↓	0.21 $\pm$ 0.05↓	1.09 $\pm$ 0.21↑*	0.98 $\pm$ 0.12↑*	0.48 $\pm$ 0.16↑
<i>Planctomycetes</i>	0.12 $\pm$ 0.02↓*	0.22 $\pm$ 0.11	0.33 $\pm$ 0.10↑	0.08 $\pm$ 0.001↓*	0.16 $\pm$ 0.07↓	0.68 $\pm$ 0.04↑*	0.54 $\pm$ 0.16↑*	0.22 $\pm$ 0.09
<i>Deferribacteres</i>	0.01 $\pm$ 0.001↓*	0.28 $\pm$ 0.12	0.05 $\pm$ 0.01↓**	0.45 $\pm$ 0.06↑*	0.61 $\pm$ 0.11↑*	0.18 $\pm$ 0L08↓	0.02 $\pm$ 0.01↓	0.41 $\pm$ 0.07↑*
<i>Candidatus Saccharibacteria</i>	0.05 $\pm$ 0.01↓**	0.46 $\pm$ 0.11	0.15 $\pm$ 0.02↓**	0.15 $\pm$ 0.02↓*	0.34 $\pm$ 0.08↓	0.05 $\pm$ 0.01↓*	0.04 $\pm$ 0.01↓*	0.07 $\pm$ 0.01↓*
<i>Verrucomicrobia</i>	0.07 $\pm$ 0.02↓	0.15 $\pm$ 0.03	0.14 $\pm$ 0.01↓	0.04 $\pm$ 0.001↓*	0.06 $\pm$ 0.01↓*	0.38 $\pm$ 0.10↑*	0.29 $\pm$ 0.07↑*	0.14 $\pm$ 0.03↓
<i>Unclassified</i>	0.05 $\pm$ 0.01↓	0.07 $\pm$ 0.01	0.07 $\pm$ 0.02↓	0.08 $\pm$ 0.023↑	0.11 $\pm$ 0.02↑	0.04 $\pm$ 0.01↓	0.13 $\pm$ 0.04↑	0.08 $\pm$ 0.01↑