**TABLE S1**. Pairwise Kruskal-Wallis test for phylogenetic diversity of treatment groups using Shannon index analysis on d 8 and 42.1

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | **Group 1** | **Group 2** | **H** | ***P-value*** | **q-value** |
| d 8 | NC | PC | 0.099 | 0.752 | 0.752 |
| PPEO | 1.334 | 0.247 | 0.743 |
| PROB | 0.275 | 0.599 | 0.752 |
| PC | PPEO | 0.099 | 0.752 | 0.752 |
| PROB | 0.397 | 0.528 | 0.752 |
| PPEO | PROB | 2.161 | 0.141 | 0.743 |
| d 42 | NC | PC | 0.656 | 0.417 | 0.501 |
|  | PPEO | 3.013 | 0.082 | 0.230 |
|  | PROB | 2.6250 | 0.105 | 0.230 |
| PC | PPEO | 2.481 | 0.115 | 0.230 |
|  | PROB | 0.705 | 0.400 | 0.501 |
| PPEO | PROB | 0.397 | 0.528 | 0.528 |

1 Treatments include: negative control (NC): corn-soybean meal basal diet; positive control (PC): NC + 20 g virginiamycin (453 g Stafac®20)/ton diet; probiotic (PROB): NC + 227 g *B. licheniformis* spores/ton diet; probiotic/prebiotic/essential oil supplement (PPEO): NC + 453 g of a supplement containing B. licheniformis spores, mannan-oligosaccharides, β-glucans (1,3 and 1,6), capsaicin and curcuma/ton diet.

2 *P*-values were calculated based on 999 permutation tests.



**(A)**

Shannon index

NC

PC

PROB

PPEO

**(B)**



NC

PC

PPEO

PROB

**FIGURE S1**. Comparison of microbial population in broiler chickens fed various dietary supplements on d 8. (**A**) Shannon’s diversity index of each treatment group. Boxplots show the quartiles, median, and extremities of the values. (**B**) 3D PCoA plots based on weighted UniFrac distance matrix (bray-curtis). Each sphere represents a sample. Treatments include: negative control (NC): corn-soybean meal basal diet; positive control (PC): NC + 20 g virginiamycin (453 g Stafac®20)/ton diet; probiotic (PROB): NC + 227 g *B. licheniformis* spores/ton diet; probiotic/prebiotic/essential oil supplement (PPEO): NC + 453 g of a supplement containing *B. licheniformis* spores, mannan-oligosaccharides, β-glucans (1,3 and 1,6), capsaicin and curcuma/ton diet.

**TABLE S2**. Pairwise PERMANOVA statistic based on unweighted UniFrac distance matrix on d 8 and 42.1

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | **Group 1** | **Group 2** | **pseudo-F** | ***P*-value**2 | **q-value** |
| d 8 | NC | PC | 0.549 | 0.887 | 0.887 |
| PPEO | 1.922 | 0.056 | 0.246 |
| PROB | 1.270 | 0.243 | 0.291 |
| PC | PPEO | 1.745 | 0.082 | 0.246 |
| PROB | 1.476 | 0.140 | 0.280 |
| PPEO | PROB | 1.205 | 0.212 | 0.291 |
| d 42 | NC | PC | 0.798 | 0.636 | 0.636 |
|  | PPEO | 1.053 | 0.374 | 0.561 |
|  | PROB | 1.374 | 0.115 | 0.230 |
| PC | PPEO | 1.742 | 0.070 | 0.230 |
|  | PROB | 1.674 | 0.096 | 0.230 |
| PPEO | PROB | 0.955 | 0.515 | 0.618 |

1 Treatments include: negative control (NC): corn-soybean meal basal diet; positive control (PC): NC + 20 g virginiamycin (453 g Stafac®20)/ton diet; probiotic (PROB): NC + 227 g *B. licheniformis* spores/ton diet; probiotic/prebiotic/essential oil supplement (PPEO): NC + 453 g of a supplement containing B. licheniformis spores, mannan-oligosaccharides, β-glucans (1,3 and 1,6), capsaicin and curcuma/ton diet.

2 *P*-values were calculated based on 999 permutation tests.

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**(A)**

Shannon index

NC

PC

PROB

PPEO



**(B)**

NC

PC

PPEO

PROB

**FIGURE S2**. Comparison of microbial population in broiler chickens fed various dietary supplements on d 42. (**A**) Shannon’s diversity index of each treatment group. Boxplots show the quartiles, median, and extremities of the values. (**B**) 3D PCoA plots based on weighted UniFrac distance matrix (bray-curtis). Each sphere represents a sample. Treatments include: negative control (NC): corn-soybean meal basal diet; positive control (PC): NC + 20 g virginiamycin (453 g Stafac®20)/ton diet; probiotic (PROB): NC + 227 g *B. licheniformis* spores/ton diet; probiotic/prebiotic/essential oil supplement (PPEO): NC + 453 g of a supplement containing *B. licheniformis* spores, mannan-oligosaccharides, β-glucans (1,3 and 1,6), capsaicin and curcuma/ton diet.