

# Gut microbiome modifications over time when removing in-feed antibiotics from the prophylaxis of post-weaning diarrhea in piglets

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## Supplementary Material

**S1 Table 1.** Average number of 16S rRNA-gene sequences (+/- standard deviation) per experimental group and timepoint

treatment	T0	T1	T2	T3
oil	55367.2+/-21348.7	81026.8+/-51291.1	69871.3+/-59729.7	110015.2+/-37496.9
traditional	46274+/-31807.1	40654.8+/-26176.7	99087.8+/-77029.3	143167+/-61104.9
white	27621.4+/-11270.9	97793.8+/-107045.6	110918.4+/-30251.9	153425+/-97899.7