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| --- | --- | --- |
| **TABLE S3** | Illumina MiSeq sequenced bacterial data based on ITS gene. | | |
| Samples | raw\_tags | clean\_tags |
| P0\_1 | 49043 | 47120 |
| P0\_2 | 47230 | 44770 |
| P0\_3 | 43117 | 41820 |
| FP\_1 | 52191 | 49008 |
| FP\_2 | 52897 | 49544 |
| FP\_3 | 48917 | 47566 |
| P1\_1 | 42116 | 40144 |
| P1\_2 | 43460 | 42024 |
| P1\_3 | 42229 | 35632 |
| P2\_1 | 46757 | 44296 |
| P2\_2 | 48881 | 47660 |
| P2\_3 | 48687 | 47597 |
| P3\_1 | 44503 | 41045 |
| P3\_2 | 42382 | 41242 |
| P3\_3 | 41754 | 40944 |
| raw\_tags, spliced from filtered and low-quality fastq data. clean\_ tags, further removing chimeras and short sequences from the splicing results. | | |