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| **TABLE S2** | Illumina MiSeq sequenced bacterial data based on 16srRNA gene. | | |
| Samples | raw\_tags | clean\_tags |
| P0\_1 | 33331 | 31258 |
| P0\_2 | 36889 | 35947 |
| P0\_3 | 38180 | 36629 |
| FP\_1 | 37550 | 36354 |
| FP\_2 | 32897 | 31989 |
| FP\_3 | 41426 | 40483 |
| P1\_1 | 41482 | 40644 |
| P1\_2 | 42520 | 41515 |
| P1\_3 | 41284 | 40114 |
| P2\_1 | 42875 | 41731 |
| P2\_2 | 41474 | 40173 |
| P2\_3 | 47105 | 45456 |
| P3\_1 | 44868 | 43992 |
| P3\_2 | 42127 | 41189 |
| P3\_3 | 49855 | 48530 |
| raw\_tags, spliced from filtered and low-quality fastq data. clean\_ tags, further removing chimeras and short sequences from the splicing results. | | |