**Table S2.** Summary of the 16S rRNA gene sequence reads during quality control assessment

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  | T1\_1 | T1\_2 | T1\_3 | T2\_1 | T2\_2 | T2\_3 |
| Raw reads | 181,851 | 169,056 | 187,258 | 181,955 | 152,505 | 189,101 |
| Filtered reads | 147,585 | 137,031 | 152,454 | 148,082 | 124,326 | 153,597 |
| Denoised reads | 141,663 | 131,755 | 146,401 | 142,533 | 119,245 | 148,239 |
| Merged reads | 117,994 | 111,858 | 119,733 | 117,589 | 98,060 | 125,090 |
| Non-chimeric reads | 104,917 | 99,843 | 104,946 | 103,882 | 86,573 | 111,146 |