**Table S2.** Analysis of sequencing data

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Samples | Raw reads | Length<18 | Length>30 | Low quality | Containing N’ reads | Clean reads |
| Bt 1 | 67,477,926 | 26,910,294 | 2,258,649 | 3,192 | 0 | 38,305,791 |
| Bt 2 | 119,103,546 | 86,371,184 | 4,001,095 | 14,526 | 0 | 28,716,741 |
| Bt 3 | 124,790,362 | 88,208,967 | 3,406,593 | 11,060 | 0 | 33,163,742 |
| CK 1 | 97,557,166 | 58,448,670 | 9,079,322 | 11,535 | 0 | 30,017,639 |
| CK 2 | 86,091,768 | 55,767,429 | 3,177,641 | 8,981 | 0 | 27,137,717 |
| CK 3 | 113,187,635 | 84,564,942 | 5,416,851 | 12,733 | 0 | 23,193,109 |