

**Table S4 Statistical information of transcriptome sequencing data**

| Sample | Clean reads | GC (%) | Q20 (%) | CycleQ20 (%) | Q30 (%) |
|--------|-------------|--------|---------|--------------|---------|
| L0     | 24,325,509  | 43     | 97      | 100          | 93      |
| L4     | 23,176,253  | 43     | 98      | 100          | 93      |
| L8     | 21,818,351  | 43     | 98      | 100          | 93      |
| L12    | 22,452,212  | 43     | 98      | 100          | 94      |
| S0     | 21,448,941  | 43     | 97      | 100          | 92      |
| S4     | 23,058,832  | 43     | 97      | 100          | 92      |
| S8     | 21,035,161  | 43     | 97      | 100          | 92      |
| S12    | 20,943,741  | 43     | 97      | 100          | 92      |