

Table 1 Summary of transcriptome data quality

Sample	Raw Reads	Clean Reads	Q20 (%)	Q30 (%)
A0h_1	49 956 086	48 345 146	97.80	93.78
A0h_2	60 683 246	59 898 532	97.87	93.80
A0h_3	61 470 694	60 069 372	97.92	93.99
A8h_1	51 587 050	50 699 510	97.92	94.03
A8h_2	52 928 692	49 932 234	97.59	93.30
A8h_3	58 296 146	55 617 338	97.99	94.16
A48h_1	56 932 668	55 630 668	97.81	93.76
A48h_2	53 354 632	49 324 524	97.97	94.06
A48h_3	51 494 542	50 737 750	98.01	94.12