

**Supplementary Table S5 General information of transcriptome data**

Sample Name	Raw Reads	Clean Reads	Clean Base(G)	Effective Rate(%)	Error Rate(%)	Q20(%)	Q30(%)	GC Content(%)
EV-1	21,258,893	20,846,291	6.25	98.06	0.03	98.09	94.19	43.61
EV-2	25,835,885	25,242,772	7.57	97.70	0.03	98.09	94.13	42.84
TX1-1	23,847,112	23,397,754	7.02	98.12	0.03	98.06	94.05	42.86
TX1-2	21,133,883	20,748,489	6.22	98.18	0.03	98.04	94.07	43.03
TX2-1	22,169,125	20,737,571	6.22	93.54	0.03	97.37	92.62	43.14
TX2-2	19,785,919	18,452,240	5.54	93.26	0.03	97.49	92.91	43.04
TX1_2-1	20,080,842	18,833,734	5.65	93.79	0.03	97.56	93.08	43.31
TX1_2-2	22,429,514	20,827,590	6.25	92.86	0.03	97.97	94.15	43.14