

Table S1 Statistical of transcriptome sequencing data

Sample	Clean reads	GC(%)	Q20(%)	Q30(%)
HA1	26036622	45.87	97.44	93.45
HA2	23408872	45.73	97.44	93.41
HA3	24055635	49.37	97.63	93.74
HT1	20789569	48.43	97.46	93.54
HT2	29494559	47.73	97.64	93.8
HT3	25650764	46.59	97.48	93.46
YA1	27679987	48.25	97.59	93.55
YA2	26977079	48.35	97.42	93.12
YA3	30308690	47.98	97.66	93.63
YT1	27135111	47.24	97.58	93.55
YT2	26614965	46.22	97.61	93.64
YT3	26248574	46.33	97.52	93.61