

Table S2. Quality control results of transcriptome sequencing data

Samples	Original reads	Filtered reads	Base error rate (%)	Q20 (%)	Q30 (%)	GC content (%)
HM_4_1_R	45228872	44702372	0.03	96.96	91.80	44.05
HM_4_2_R	44003114	43508386	0.03	96.76	91.33	44.09
HM_4_3_R	46176524	45742780	0.03	96.76	91.36	44.23
HM_6_1_R	47451968	46878202	0.03	96.80	91.42	44.70
HM_6_2_R	46667688	46195772	0.03	97.03	91.88	44.42
HM_6_3_R	42923478	42620382	0.03	97.13	92.05	44.38
HM_7_1_R	47229012	46524078	0.03	96.69	91.42	42.93
HM_7_2_R	44418972	43939798	0.03	96.52	90.82	44.36
HM_7_3_R	47223980	46722434	0.03	96.89	91.64	44.44
JJX_4_1_R	47407522	46841026	0.03	96.83	91.42	43.65
JJX_4_2_R	49287808	48811472	0.03	96.98	91.78	43.76
JJX_4_3_R	46315532	45739908	0.03	97.12	92.03	43.77
JJX_6_1_R	48070832	47427522	0.03	97.13	92.06	43.86
JJX_6_2_R	45821558	45123820	0.03	96.98	91.83	44.23
JJX_6_3_R	43525112	42619824	0.03	96.88	91.57	44.11
JJX_7_1_R	41678748	40973636	0.03	96.98	91.77	44.08
JJX_7_2_R	45631398	44725708	0.03	96.85	91.51	44.04
JJX_7_3_R	44843856	44317792	0.03	96.98	91.76	44.00

Note: HM_4, HM_6, and HM_7 represent 78, 92, and 99 DAB ‘Huami’ wampee fruits, respectively. JJX_4, JJX_6, and JJX_7 represent 78, 92, and 99 DAB ‘Jinjixin’ wampee fruits, respectively.