

Table S1 quality analysis of transcriptome sequencing data

sample name	Raw reads	Clean reads	Error rate %	Q20 %	Q30 %	GC content %	Contrast ratio%
HJH Albedo 1	41725292	41301096	0.0257	97.8	93.38	45.24	88.87
HJH Albedo 2	50405050	49618582	0.0256	97.85	93.52	45.1	90.96
HJH Albedo 3	44951404	44363436	0.0262	97.59	92.9	44.9	92.05
R-HJH Albedo 1	52446264	51948746	0.0257	97.79	93.38	44.52	93.12
R-HJH Albedo 2	49635476	49171630	0.026	97.7	93.14	44.83	92.7
R-HJH Albedo 3	47521580	47099778	0.0257	97.79	93.36	44.93	90.69