

Changes have been marked in red

Supplemental Data

Supplementary Table S1. Assemble and compare transcriptome data of strawberry

Sample	Total Raw Reads (M)	Total Clean Reads (M)	Total Clean Base(Gb)	Q30 (%)	Total Mapping(%)
0d_1	43.82	42.63	6.39	92.48	89.00
0d_2	43.82	42.45	6.37	92.72	89.18
0d_3	43.82	42.52	6.38	92.68	89.12
2d_1	43.82	40.32	6.05	92.47	89.19
2d_2	43.82	42.54	6.38	92.72	89.00
2d_3	43.82	41.47	6.24	92.61	89.13
4d_1	43.82	42.30	6.34	92.72	89.04
4d_2	43.82	42.63	6.39	92.79	88.96
4d_3	43.82	42.58	6.36	92.74	88.79
6d_1	43.82	42.41	6.36	92.65	89.01
6d_2	43.82	42.62	6.39	92.76	89.28
6d_3	43.82	42.53	6.37	92.65	89.13
8d_1	43.82	42.25	6.34	92.84	89.01
8d_2	43.82	42.39	6.36	92.62	88.97
8d_3	43.81	42.32	6.33	92.79	88.85