

Table S2. Data of sequencing result

Sample	Raw_reads	Raw_bases	Clean_reads	Clean_bases	Error_rate	Q20	Q30	GC_pct
black_1	41322628	6.2G	41322628	6.2G	0.03	97.95	93.88	45.09
black_2	43382688	6.51G	43382688	6.51G	0.03	97.58	93.13	44.91
black_3	42742170	6.41G	42742170	6.41G	0.03	97.97	93.91	44.91
yellow_1	41649428	6.25G	41649428	6.25G	0.03	97.17	92.34	40.91
yellow_2	41901754	6.29G	41901754	6.29G	0.03	97.63	93.24	42.26
yellow_3	38280086	5.74G	38280086	5.74G	0.03	97.38	92.71	42.22
Dark red_1	39924190	5.99G	39924190	5.99G	0.03	97.49	92.89	42
Dark red_2	39727362	5.96G	39727362	5.96G	0.03	97.66	93.34	41.76
Dark red_3	41854842	6.28G	41854842	6.28G	0.03	97.79	93.56	41.73
yellow_with_s_1	41677196	6.25G	41677196	6.25G	0.03	97.88	93.81	41.71
yellow_with_s_2	40632966	6.09G	40632966	6.09G	0.03	97.72	93.45	41.82
yellow_with_s_3	41339002	6.2G	41339002	6.2G	0.03	97.47	92.86	41.91