

Supplementary Table 4. The statistics of *Camellia chekiangoleosa* transcriptome. The transcripts and unigenes are obtained by using the Trinity v2.11.0 software. Sequences less than 200 bp are removed.

	All	GC%	Min Length (bp)	Median Length (bp)	Max Length (bp)	Total Assembled Bases (bp)	N50 (bp)
Transcript	206367	40.79	201	684	13728	202547343	1492
Unigene	40042	40.85	201	835	13728	44816160	1676