

Table S7. The statistic of RNA-seq data.

Sample	Reads count	Base (bp)	GC(%)
sepal_1	41247514	12374254200	45
sepal_2	41494983	12448494900	46
Petal_1	41987624	12596287200	45
Petal_2	38146914	11444074200	46
stigma	43872373	13161711900	45
carpel	42831599	12849479700	44
ovary	40882800	12264840000	44
anther	40713592	12214077600	46
filament	45216626	13564987800	45
receptacle	44267177	13280153100	45
pedicel	43651659	13095497700	45