

Table S1. Primers used in this study.

Genes		Sequence(5' to 3')	Fragment size(bp)
MYB80	Forward primer	CACTCTGTCGTGGAAAC	154
	Reverse primer	TAAGGTGGGAGAAAGGCT	
AMS	Forward primer	CAAACGGCTGGGACTACT	225
	Reverse primer	CATGTATCCCGATTGAGG	
GSL5	Forward primer	GGCATTAGTAAGGCATCC	138
	Reverse primer	CTGGTTCAGTCCAACATCA	
TDF1	Forward primer	GTCAGGACCCATCACAAAC	194
	Reverse primer	CAAGGAAATCACTCCACTC	
SERK2-like	Forward primer	TCTACGCCTTGGTGTC	143
	Reverse primer	AGCAGCCTCCTGTTTGA	
AGAMOUS	Forward primer	TCATCTCCTCCTCCGTT	238
	Reverse primer	CCTCTTCAGCAGTCATT	
LLA-115	Forward primer	GGGAAATGACGACACCTT	150
	Reverse primer	CACCACTGAATCCAAACCT	
NOZZLE	Forward primer	AGACGGAGGTGACGGAGTA	117
	Reverse primer	CAACAAAATGAGGAGACGGT	
GLOX1	Forward primer	ATTCAGTCTGGCGGCTAC	194
	Reverse primer	GATTGGGGATGAACTCGTA	
ARF17	Forward primer	TTTCCCAAAGGTGCTAAC	118
	Reverse primer	CTGGAATGGCAGATAAAGAG	
MYB24	Forward primer	CAACCACCTCTCCTTCA	135
	Reverse primer	CTCCGCTTCTTCTCCACC	
FPA	Forward primer	AGCCCTTAGCCATTCAC	178
	Reverse primer	TCTGCTTCAGCCTTCC	
CYP704B1	Forward primer	CAAACGCCACTGTCACGC	233
	Reverse primer	GTCTTGTTCCTGCCAC	
chalcone synthase (CHS)	Forward primer	GGGCTAACATTCCACCTA	116
	Reverse primer	CAGAAAATCGAGTCCAGTC	
chalcone isomerase (CHI)	Forward primer	TTTCCTGCTGCCACTCC	124
	Reverse primer	GCTCCCACCCAAATACCA	
cysteine proteininase (CYS)	Forward primer	CGTTGAAGGGAAATCATAAG	162
	Reverse primer	CAGTTGCCTGTTGTAAGC	
18S	Forward primer	AGTTGGTGGAGCGATTGTCT	
	Reverse primer	CCTGTTATTGCCTCAAACCTTCC	

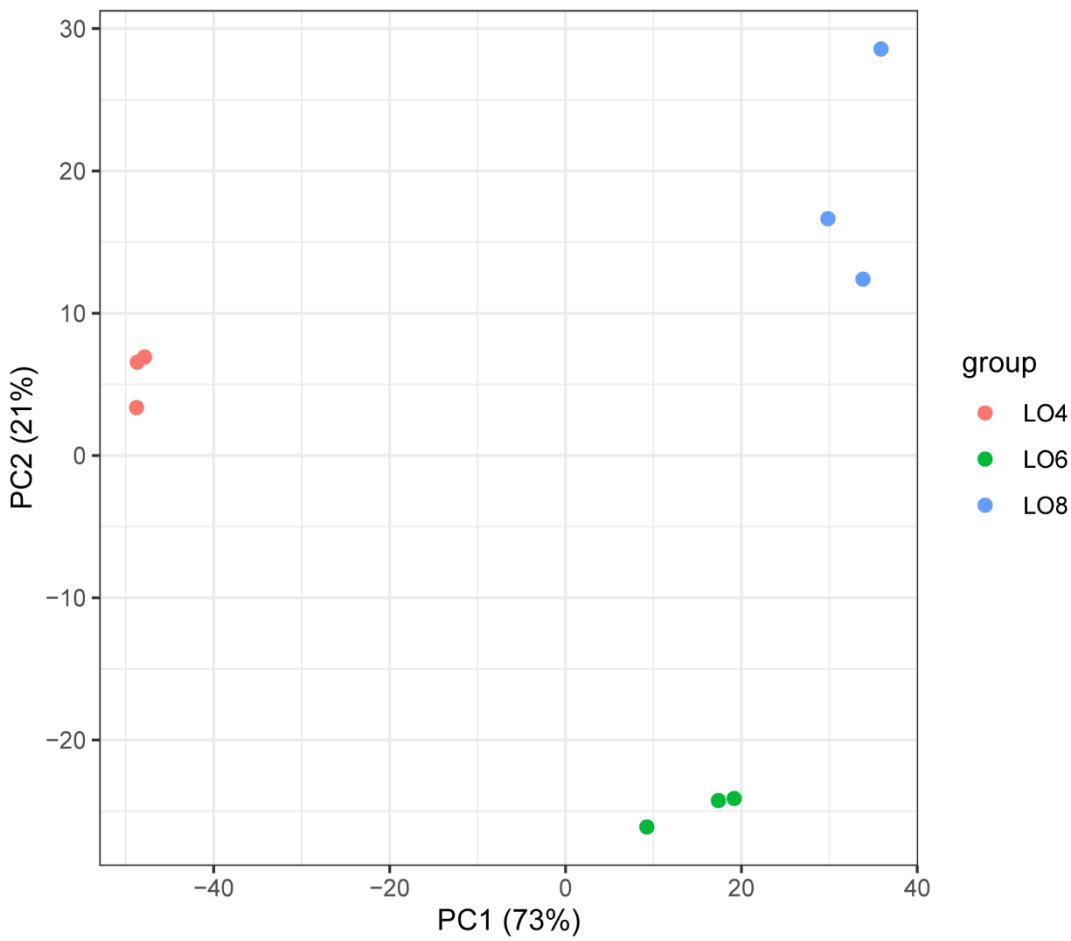


Figure S1. Biological replicates of three anther stages showed by the first two PCs which explained over 94% of variance among samples.

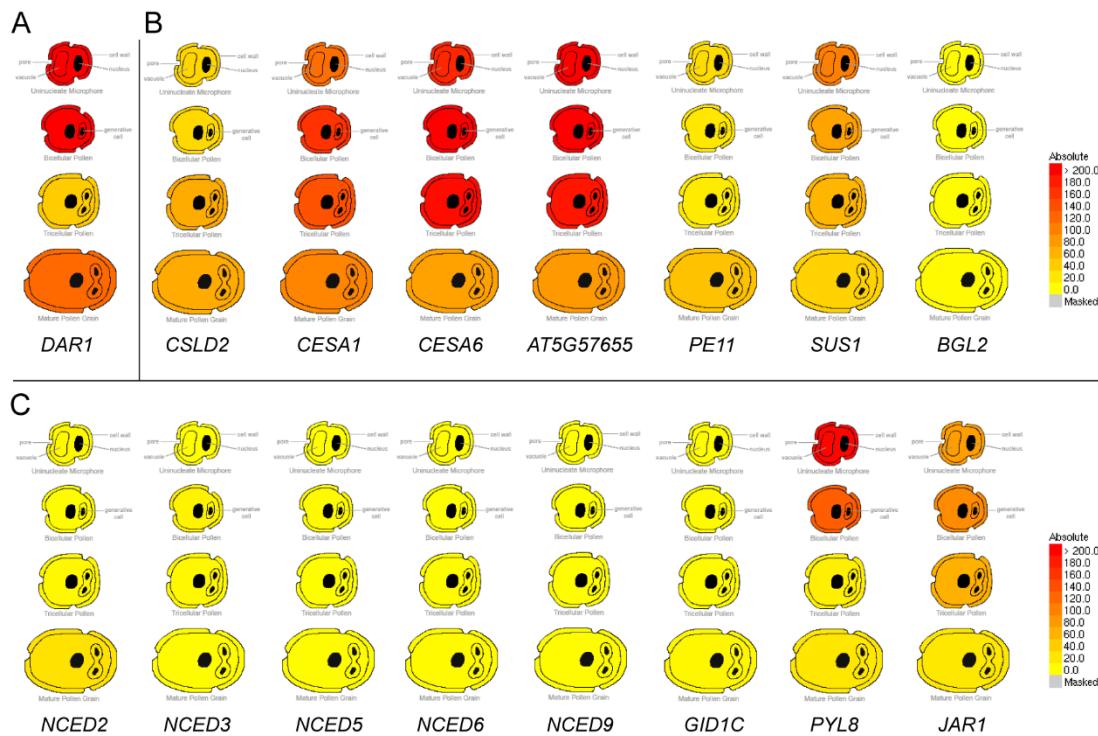


Figure S2. The 'eFP-seq' browser of genes related to the tapetum degradation and microspore development. The color scale from yellow to red represents expression levels from low to high.