

Additional file 1 Detailed information of selected datasets

Data type	Tissue type	Accession number	Abbreviation used in the present analysis
Genomic	Leaf	SRR26668329	RADSeq_L1
		SRR26668213	RADSeq_L2
		SRR26668372	RADSeq_L3
		SRR26668382	RADSeq_L4
		SRR26668312	RADSeq_L5
		SRR26668309	RADSeq_L6
		SRR26668303	RADSeq_L7
		SRR26668248	RADSeq_L8
		SRR26668247	RADSeq_L9
		SRR26668345	RADSeq_L10
		SRR26668340	RADSeq_L11
		SRR26668278	RADSeq_L12
		SRR26668264	RADSeq_L13
		SRR26668262	RADSeq_L14
Transcriptomic	Root	SRR8816492	RNASeq_R1
		SRR8816486	RNASeq_R2
	Stem	SRR8816469	RNASeq_S1
		SRR8816470	RNASeq_S2
	Flower	SRR8816488	RNASeq_F1
		SRR8816489	RNASeq_F2

Leaf	SRR8816484	RNASeq_L1
	SRR8816485	RNASeq_L2
2 MAP*	SRR8816471	RNASeq_2MAP1
	SRR8816472	RNASeq_2MAP2
4 MAP*	SRR8816474	RNASeq_4MAP1
	SRR8816475	RNASeq_4MAP2
6 MAP*	SRR8816477	RNASeq_6MAP1
	SRR8816478	RNASeq_6MAP2
8 MAP*	SRR8816480	RNASeq_8MAP1
	SRR8816482	RNASeq_8MAP2

*MAP: months after pollination