

Table S7. Overlap between tissue in expression profiles gene ontology term enrichment for Biological processes (A) and Molecular functions (B).

A

Categories	Biological processes				
	F vs L	F vs R	L vs R	F vs L vs R	Single
Reverse	0	22	11	0	67
Transgressive	3	14	10	4	69
Legacy	2	9	7	2	80
Dominance CG	8	8	3	0	81
Comp. drift	5	7	2	0	86
Intermediate	3	2	5	2	88
Dominance CO	3	6	0	0	91
Average	3	10	5	1	80

B

Categories	Molecular functions				
	F vs L	F vs R	L vs R	F vs L vs R	Single
Transgressive	2	15	4	7	72
Reverse	0	10	10	5	75
Intermediate	2	9	8	3	78
Legacy	5	11	3	2	79
Dominance CG	5	7	8	0	80
Comp. Drift	4	7	6	2	81
Dominance CO	4	5	1	4	86
Average	3	9	6	3	79