

Trinity transcrip name	Annotation from the SwissProt database	BR vs 2D4W		BR vs 7D4W		2D4W vs 7D4W	
		logFC	FDR	logFC	FDR	logFC	FDR
ABA biosynthesis							
TR8162 c0_g1_i1	ABI five-binding protein 3 [<i>Arabidopsis thaliana</i>]	-0.910	7.04E-12	2.095	6.10E-22	-3.054	1.12E-40
TR8743 c0_g2_i1	9-cis-epoxycarotenoid dioxygenase NCED1 [<i>Phaseolus vulgaris</i>]	-6.157	1.69E-28	0.082	1.00	-6.307	3.71E-30
TR1393 c0_g1_i1	ABA-HYPERSENSITIVE GERMINATION 3; Protein phosphatase 2C A [<i>Arabidopsis thaliana</i>]	0.525	0.0310	3.136	1.16E-56	-2.664	6.35E-30
TR8214 c0_g1_i1	ABI five-binding protein 2 [<i>Arabidopsis thaliana</i>]	1.208	2.08E-15	3.405	3.06E-100	-2.239	2.57E-25
TR13511 c0_g2_i1	Abscisic acid 8'-hydroxylase 3; Cytochrome P450 707A3 [<i>Arabidopsis thaliana</i>]	-4.664	4.69E-05	-0.919	0.889	-3.911	0.0027
GA biosynthesis							
TR9163 c0_g2_i1	Gibberellin 2-beta-hydroxylase 8 [<i>Arabidopsis thaliana</i>]	-0.725	0.175	0.490	0.474	-1.274	0.0034
MADS box							
TR15408 c4_g2_i5	MADS-box protein JOINTLESS, LeMADS [<i>Solanum lycopersicum</i>]	3.056	4.23E-88	3.733	1.18E-152	-0.732	2.78E-02
Stress respond							
TR24454 c0_g1_i1	RecName: Full=EID1-like F-box protein 3 [<i>Arabidopsis thaliana</i>]	-3.474	1.58E-41	1.878	0.0005	-5.413	4.92E-71
TR10580 c0_g2_i1	RecName: Full=Dehydration-responsive element-binding protein 2C [<i>Arabidopsis thaliana</i>]	0.472	1.63E-05	3.826	5.90E-91	-3.411	1.10E-54
TR10071 c0_g2_i1	Late embryogenesis abundant protein D-34 [<i>Gossypium hirsutum</i>]	3.771	5.18E-109	5.979	3.55E-153	-2.237	0.0002