

**Table S1. List of Primers used in this study**

Experiment	Sequence			
	Primer name	Forward	Primer name	Reverse
Construction	BamHI-GRf	ATGGGATCCCCTGAAGCTCGAAAAACAAAG	KpnI-GRr	ATGGGTACCTCATTTCGATGAAACAGAAG
	SacII-WOX1f	ATGCCGCGGATGTGGACGATGGGTTACAAC	WOX1BlcL1_1	AGTGGATCCTGCTCCTGCGTTCCTCAATGGCAGAA
	SacII-G3GFPf	ATGCCGCGGATGAGTAAAGGAGAAGAACTT	BamHI-G3GFPPr	AGTGGATCCTTTGTATAGTTCATCCATGCC
RT-qPCR	ACT2f	TCCCTCAGCACATTCCAGCAGAT	ACT2r	AACGATTCCTGGACCTGCCTCATC
	CUC2f	TGCCATCGCAGAAGTTGATC	CUC2r	TTCTCTCCCATCTTAGCTCTCCCG
	CUC3f	TCTGGTAGCTCCCAAGGTTACAG	CUC3r	TGGTCTCTCCGGCATTACATTC
	MPf	TACGCCACTTAGAACCAGGCCATC	MPr	CGGGAAACGAAGGTACATGGGAAC
	PIN1f	ACAAAACGACGCAGGCTAAG	PIN1r	AGCTGGCATTTCATGTTCC

**Table S2. List of log<sub>2</sub> expression level of DEX-up-IAA-up genes (>1.5-fold)**

ProbeName	GeneName	AGI code	EtOH_1	EtOH_2	DEX+EtOH_1	DEX+EtOH_2	EtOH+IAA_1	EtOH+IAA_2	DEX+IAA_1	DEX+IAA_2
A_84_P789756	AA042562		-4.89	-4.32	-3.27	-3.32	-3.41	-3.22	-2.93	-2.66
A_84_P13037	AAP7	AT5G23810	0.27	-0.11	2.33	1.48	1.25	1.56	3.12	2.70
A_84_P767740	AAP7	AT5G23810	-3.66	-3.67	-0.85	-2.15	-2.41	-2.23	-0.53	-0.86
A_84_P20410	ADS1	AT1G06080	2.61	2.91	6.29	5.96	4.47	5.27	6.99	7.42
A_84_P820492	AFP3	AT3G29575	-0.34	-0.19	1.14	1.21	2.66	2.60	2.07	1.92
A_84_P203128	AFP3	AT3G29575	-0.47	-0.67	1.09	1.09	2.52	2.52	1.96	1.82
A_84_P286390	AOC3	AT3G25780	-0.83	-0.67	0.39	-0.01	0.60	0.33	1.34	0.96
A_84_P17366	AT1G01250	AT1G01250	-2.78	-2.28	-0.11	0.29	-1.81	-1.18	-0.10	0.35
A_84_P22852	AT1G01520	AT1G01520	-2.65	-3.32	-1.17	-2.31	-1.31	-1.88	-1.31	-2.10
A_84_P838948	AT1G01520	AT1G01520	-4.01	-4.66	-3.03	-4.45	-2.99	-3.76	-3.09	-4.18
A_84_P842621	AT1G02670	AT1G02670	-2.31	-1.90	-1.70	-1.09	-1.86	-1.48	-1.19	-0.74
A_84_P606594	AT1G02670	AT1G02670	-4.85	-3.94	-3.40	-3.31	-3.41	-3.31	-3.16	-2.71
A_84_P16098	AT1G03730	AT1G03730	2.81	2.94	3.30	2.77	3.62	3.04	3.32	2.91
A_84_P851101	AT1G03730	AT1G03730	-4.35	-3.87	-2.92	-3.22	-3.42	-3.06	-2.81	-2.67
A_84_P15190	AT1G04330	AT1G04330	-3.60	-3.07	-2.26	-1.74	-1.76	-1.85	-1.07	-0.79
A_84_P522242	AT1G07985	AT1G07985	-2.47	-3.41	-1.21	-1.89	-0.72	-1.60	-0.31	-1.52
A_84_P20899	AT1G10640	AT1G10640	2.62	2.94	4.16	4.42	3.47	4.01	4.05	4.24
A_84_P555923	AT1G11125	AT1G11120	0.82	1.29	2.00	2.34	2.03	2.35	2.70	2.92
A_84_P129976	AT1G13360	AT1G13360	2.89	2.82	3.56	3.50	3.53	3.54	4.01	3.96
A_84_P860600	AT1G13360	AT1G13360	-0.30	-0.38	0.52	0.39	0.40	0.42	1.02	0.70
A_84_P12487	AT1G14390	AT1G14390	-0.94	-0.69	0.07	0.46	0.62	0.71	1.12	1.23
A_84_P20976	AT1G19210	AT1G19210	-4.74	-4.13	-2.72	-3.32	-3.65	-2.90	-3.14	-1.69
A_84_P21770	AT1G23060	AT1G23060	-1.48	-1.55	-0.44	-0.52	0.94	0.84	1.60	1.34
A_84_P266910	AT1G47655	AT1G47655	-0.81	-0.49	0.26	0.77	1.30	1.49	1.96	2.14
A_84_P15213	AT1G47670	AT1G47670	3.28	3.62	3.96	4.43	4.03	4.34	4.72	5.01
A_84_P19573	AT1G48100	AT1G48100	2.16	1.96	3.09	3.01	2.83	2.71	2.81	2.53
A_84_P23351	AT1G48180	AT1G48180	-4.83	-4.62	-3.44	-3.11	-3.33	-3.64	-4.13	-3.08
A_84_P168923	AT1G48330	AT1G48330	0.66	1.14	1.38	2.32	1.39	2.13	2.19	2.91
A_84_P23648	AT1G51170	AT1G51170	1.08	1.43	1.68	2.19	2.34	2.57	2.43	2.50
A_84_P15194	AT1G54020	AT1G54020	-3.15	-3.16	1.07	2.08	-1.29	-1.24	1.67	1.56
A_84_P503731	AT1G54575	AT1G54575	-0.26	-1.55	0.61	0.17	0.43	-0.43	0.60	-0.11
A_84_P818130	AT1G62610	AT1G62610	-0.05	0.01	0.40	0.75	0.29	0.48	0.54	0.80
A_84_P596994	AT1G62610	AT1G62610	-3.72	-3.70	-3.05	-2.61	-3.11	-3.03	-3.02	-2.66
A_84_P14874	AT1G62770	AT1G62770	-1.33	-1.18	-0.44	-0.03	0.83	1.61	1.36	2.19
A_84_P12439	AT1G68110	AT1G68110	-3.89	-3.60	-2.43	-2.84	-3.06	-3.00	-3.10	-2.59
A_84_P19840	AT1G71691	AT1G71691	-1.50	-0.78	-0.71	0.04	-0.14	-0.14	-0.47	0.23
A_84_P750168	AT1G75060	AT1G75060	0.74	0.99	1.12	1.65	1.29	1.72	1.61	1.94
A_84_P524195	AT1G75060	AT1G75060	-0.22	0.05	0.44	0.92	0.40	0.79	0.81	1.15
A_84_P18113	AT1G76640	AT1G76640	-3.73	-3.74	-1.24	-0.82	-1.17	-0.91	-0.29	0.04
A_84_P73664	AT1G77855	AT1G77855	-1.02	-0.78	1.01	1.10	0.10	0.23	1.69	1.59
A_84_P281740	AT1G78070	AT1G78070	1.95	1.92	2.73	2.46	2.92	2.60	3.11	2.57
A_84_P785283	AT1G78070	AT1G78070	0.76	0.57	1.57	1.34	1.62	1.39	1.97	1.35
A_84_P147538	AT1G80865	AT1G80865	-4.49	-4.13	-2.44	-1.60	-3.70	-3.13	-2.26	-1.86
A_84_P246685	AT2G07360	AT2G07360	5.04	5.06	5.05	5.21	5.02	5.12	5.08	5.22
A_84_P819612	AT2G07360	AT2G07360	3.97	3.98	4.04	4.25	4.00	4.07	4.05	4.25
A_84_P803354	AT2G07360	AT2G07360	-4.12	-4.27	-3.51	-2.97	-3.38	-3.59	-3.10	-3.30
A_84_P755231	AT2G14270	AT2G14270	0.73	1.07	1.60	1.76	1.35	1.70	1.95	2.35
A_84_P758125	AT2G16668	AT2G16668	-4.68	-4.71	-3.68	-3.75	-3.81	-3.97	-3.76	-3.66
A_84_P98146	AT2G17500	AT2G17500	2.14	1.20	3.00	1.97	3.49	2.95	3.66	2.94
A_84_P816631	AT2G17500	AT2G17500	-0.54	-1.45	0.72	-0.61	0.86	0.22	1.12	0.47
A_84_P555310	AT2G36650	AT2G36650	-2.87	-2.71	-1.86	-2.02	-1.95	-2.11	-2.00	-2.22
A_84_P13527	AT2G37030	AT2G37030	-1.60	-2.43	-0.20	-0.89	0.57	0.96	3.22	2.85
A_84_P15421	AT2G37760	AT2G37760	1.16	-0.22	1.76	1.05	2.05	0.86	1.91	1.15
A_84_P820187	AT2G37760	AT2G37760	-1.15	-2.49	-0.38	-1.10	-0.36	-1.37	-0.19	-0.96
A_84_P586474	AT2G39855	AT2G39855	-3.60	-3.40	-1.19	-1.30	-1.56	-1.12	0.54	0.56
A_84_P23844	AT2G43870	AT2G43870	-2.11	-1.95	-0.60	-0.34	0.01	0.08	1.40	1.79
A_84_P506584	AT2G46375	AT2G46375	-3.70	-3.81	-2.66	-2.75	-2.95	-3.13	-2.47	-3.72
A_84_P216308	AT2G47270	AT2G47270	-4.79	-4.02	-1.84	-2.42	-1.18	-1.19	-0.36	-0.11
A_84_P125091	AT3G04860	AT3G04860	0.96	1.10	1.73	2.25	1.69	1.85	2.20	2.23
A_84_P242043	AT3G19500	AT3G19500	-4.87	-3.70	-3.59	-2.74	-3.88	-3.09	-3.76	-2.90
A_84_P559854	AT3G20830	AT3G20830	-0.50	-0.65	-0.07	0.29	0.37	0.33	0.79	0.54
A_84_P841540	AT3G20830	AT3G20830	-0.69	-0.63	-0.03	0.17	0.35	0.36	0.78	0.55
A_84_P23101	AT3G21890	AT3G21890	-1.01	-2.03	0.10	-0.95	0.03	-0.55	0.11	-0.96
A_84_P507571	AT3G25130	AT3G25130	-0.95	-0.49	-0.06	0.37	-0.15	0.14	0.56	0.81
A_84_P117052	AT3G25640	AT3G25640	0.17	0.67	1.02	1.62	1.07	1.49	1.72	2.35
A_84_P762469	AT3G26742	AT3G26742	-3.90	-3.47	-2.48	-2.61	-2.77	-2.31	-1.85	-1.66
A_84_P787999	AT3G29370	AT3G29370	-0.35	-0.47	1.00	0.77	3.69	3.65	4.13	4.17
A_84_P144819	AT3G29370	AT3G29370	-1.74	-1.81	-0.23	-0.44	2.49	2.42	2.90	2.92
A_84_P286640	AT3G48510	AT3G48510	-2.60	-2.97	-1.98	-1.84	-1.51	-1.42	-1.59	-1.92
A_84_P20327	AT3G50400	AT3G50400	-4.10	-3.68	-2.90	-2.71	-3.29	-2.58	-2.28	-1.95
A_84_P592005	AT3G57380	AT3G57380	-3.52	-3.94	-1.68	-2.20	-2.32	-3.11	-1.78	-2.60
A_84_P869477	AT3G60520	AT3G60520	1.33	1.10	2.87	2.60	2.99	3.14	3.96	3.96

A_84_P304930	AT3G60520	AT3G60520	0.79	0.60	2.56	2.27	2.60	2.64	3.45	3.47
A_84_P17537	AT3G61680	AT3G61680	-2.82	-2.65	-0.68	-0.84	-0.50	-0.01	0.98	1.10
A_84_P764577	AT4G14819	AT4G14819	-1.96	-1.87	-0.97	-0.61	2.04	1.96	2.26	2.14
A_84_P502380	AT4G19970	AT4G19970	-4.06	-3.77	-2.27	-1.91	-3.42	-2.82	-2.04	-1.79
A_84_P765231	AT4G20320	AT4G20320	0.85	1.47	2.16	2.40	1.58	2.02	2.46	2.97
A_84_P831942	AT4G20320	AT4G20320	-2.44	-2.02	-0.90	-0.67	-1.80	-1.24	-0.86	-0.22
A_84_P833441	AT4G20320	AT4G20320	-4.10	-3.80	-2.61	-2.37	-3.38	-2.67	-2.30	-1.92
A_84_P231229	AT4G21930	AT4G21930	-2.33	-3.10	-0.74	-1.17	-1.59	-1.85	-1.97	-2.07
A_84_P522811	AT4G23720	AT4G23720	-1.59	-1.08	-0.56	-0.31	-0.20	0.09	0.67	0.80
A_84_P18585	AT4G28650	AT4G28650	-1.76	-1.46	-0.49	-0.11	0.09	0.26	0.09	0.45
A_84_P825877	AT4G28650	AT4G28650	-4.42	-3.98	-2.61	-2.31	-2.08	-2.09	-2.20	-1.98
A_84_P593014	AT4G31110	AT4G31110	-2.65	-2.17	-2.11	-2.03	-2.16	-1.81	-2.09	-2.03
A_84_P789549	AT4G31110	AT4G31110	-4.47	-4.18	-3.86	-3.28	-3.54	-3.26	-3.14	-3.11
A_84_P15762	AT4G31910	AT4G31910	0.46	0.80	1.26	1.96	1.92	2.45	2.68	2.99
A_84_P583210	AT4G33550	AT4G33550	-2.42	-2.14	-0.68	0.05	-1.04	-0.46	-1.02	-0.37
A_84_P549446	AT4G35720	AT4G35720	0.15	0.38	1.41	1.24	2.20	2.71	2.67	3.07
A_84_P233347	AT4G36010	AT4G36010	1.76	2.09	2.47	3.15	2.56	2.88	2.71	2.86
A_84_P117782	AT4G37690	AT4G37690	-4.22	-3.98	-3.38	-3.00	-3.50	-3.35	-3.61	-2.80
A_84_P740964	AT5G08760	AT5G08760	1.14	0.57	1.95	1.45	1.74	1.80	2.69	2.11
A_84_P843324	AT5G08760	AT5G08760	-0.73	-1.07	0.06	-0.34	0.04	-0.09	0.77	0.16
A_84_P570789	AT5G11360	AT5G11360	-4.77	-4.27	-3.13	-3.15	-3.65	-3.20	-2.78	-2.71
A_84_P542432	AT5G20810	AT5G20810	-3.79	-4.20	-3.03	-2.77	-3.17	-2.79	-3.22	-2.50
A_84_P592224	AT5G23100	AT5G23100	-0.18	0.02	0.95	1.54	0.58	0.62	1.29	1.78
A_84_P199084	AT5G38200	AT5G38200	1.90	1.92	3.54	3.21	3.38	3.51	4.01	4.26
A_84_P10228	AT5G38820	AT5G38820	-3.96	-3.44	-3.17	-2.48	-2.91	-2.37	-2.83	-2.08
A_84_P574406	AT5G39240	AT5G39240	0.22	1.32	2.13	3.21	2.70	3.03	3.47	3.77
A_84_P595814	AT5G40800	AT5G40800	-3.75	-4.14	-2.42	-3.00	-1.72	-1.77	-2.76	-2.82
A_84_P544283	AT5G48175	AT5G48175	-2.47	-2.66	-0.58	-1.68	-1.43	-1.08	-0.09	-0.75
A_84_P586962	AT5G56880	AT5G56880	-2.96	-2.50	-1.22	-0.74	-1.83	-0.91	-0.79	-0.20
A_84_P840907	AT5G56880	AT5G56880	-4.57	-3.62	-2.90	-2.57	-3.52	-2.60	-2.47	-1.99
A_84_P825325	AT5G57340	AT5G57340	0.28	0.39	0.88	1.13	1.10	1.25	1.05	1.34
A_84_P767982	AT5G57340	AT5G57340	-2.44	-2.32	-1.30	-1.19	-1.61	-1.16	-1.23	-0.79
A_84_P559359	AT5G57785	AT5G57785	0.43	0.77	1.46	1.74	1.92	2.28	2.04	2.49
A_84_P553153	AT5G57790	AT5G57790	-1.54	-1.17	-0.27	0.11	0.00	0.42	0.56	0.88
A_84_P134445	AT5G60760	AT5G60760	-2.29	-2.01	-1.02	-0.64	-1.11	-1.30	-0.37	-0.07
A_84_P13151	AT5G61820	AT5G61820	3.51	2.62	4.19	2.93	4.20	3.20	4.03	2.58
A_84_P856918	AT5G61820	AT5G61820	-1.12	-2.13	0.01	-1.52	-0.29	-1.36	-0.25	-1.84
A_84_P793448	AT5G61820	AT5G61820	-2.87	-3.46	-2.25	-3.02	-2.30	-3.14	-2.12	-3.53
A_84_P23571	AT5G66080	AT5G66080	-1.13	-0.82	0.78	0.91	1.01	1.45	2.36	2.48
A_84_P597692	AT5G66450	AT5G66450	2.75	2.84	2.54	2.98	2.66	2.91	2.64	2.90
A_84_P850105	AT5G66450	AT5G66455	-2.56	-2.22	-1.69	-1.32	-1.97	-1.49	-1.73	-1.13
A_84_P792311	AT5G66450	AT5G66450	-1.46	-1.25	-0.85	-0.37	-1.21	-0.70	-0.48	-0.06
A_84_P21681	AT5G66460	AT5G66460	-0.16	-0.17	0.68	0.77	1.53	1.53	1.27	1.30
A_84_P223759	AT5G66580	AT5G66580	0.59	0.86	1.30	2.39	3.37	3.49	3.78	3.63
A_84_P21683	AT5G66940	AT5G66940	1.11	1.86	3.00	3.91	3.70	3.72	4.53	4.63
A_84_P21341	ATPMPEPCRB	AT4G02330	2.68	2.65	3.68	3.51	3.48	3.23	4.09	3.56
A_84_P751391	AY087114	AT1G21525	-3.85	-3.64	1.01	-0.53	-2.43	-1.72	1.89	1.15
A_84_P582920	BAP2	AT2G45760	-4.64	-4.82	-2.05	-2.64	-3.81	-4.14	-2.67	-3.80
A_84_P14435	BAS1	AT2G26710	-1.45	-0.95	-0.03	0.99	0.80	0.98	2.03	2.02
A_84_P23646	BCA3	AT1G23730	-0.56	-0.94	0.58	0.30	1.01	0.88	1.79	1.52
A_84_P850621	BIM1	AT5G08130	3.57	3.60	3.87	4.23	4.12	4.31	4.21	4.50
A_84_P19621	BIM1	AT5G08130	2.73	2.67	2.85	3.34	3.19	3.50	3.42	3.72
A_84_P810630	BIM1	AT5G08130	-3.05	-2.43	-2.25	-1.78	-1.83	-1.51	-1.88	-1.69
A_84_P795619	BIM1	AT5G08130	-0.32	-0.41	-0.32	0.12	0.09	0.30	0.23	0.31
A_84_P851509	BP670694		-3.18	-3.47	-2.13	-2.34	-2.53	-2.60	-2.73	-2.45
A_84_P819429	BPEp	AT1G59640	-1.53	-0.83	-0.17	0.10	-0.82	-0.68	-0.25	0.00
A_84_P562234	BPEp	AT1G59640	-2.12	-1.71	-0.65	-0.57	-1.27	-1.11	-0.92	-0.64
A_84_P16727	BRU6	AT4G37390	-0.32	0.22	0.88	1.97	5.18	4.82	5.99	5.54
A_84_P769292	BX833772	AT5G03552	-0.92	-0.85	0.11	0.14	-0.30	-0.05	0.23	0.35
A_84_P10585	CDS1	AT1G62430	1.84	2.27	2.45	3.01	2.41	2.80	2.79	3.06
A_84_P821514	CDS1	AT1G62430	-1.82	-1.36	-0.85	-0.49	-1.08	-0.71	-0.78	-0.57
A_84_P20987	chr31	AT1G05490	-3.81	-3.14	-2.38	-1.72	-2.74	-2.09	-2.12	-1.39
A_84_P837470	chr31	AT1G05490	-3.99	-2.96	-2.64	-1.94	-2.77	-2.36	-2.06	-1.59
A_84_P126931	CLE16	AT2G01505	-0.37	0.16	0.39	0.93	-0.37	0.26	0.30	0.80
A_84_P789435	CLE16	AT2G01505	-3.69	-3.59	-2.50	-2.28	-2.98	-2.75	-2.61	-2.41
A_84_P14345	CPK29	AT1G76040	1.57	1.36	3.33	2.35	2.01	1.79	3.72	2.73
A_84_P805748	CPK29	AT1G76040	-2.12	-2.25	-0.30	-1.40	-1.49	-1.59	0.18	-0.89
A_84_P18479	CYCP3;2	AT3G60550	-2.82	-3.23	-1.63	-2.07	-1.65	-2.11	-1.37	-1.69
A_84_P286120	CYP707A1	AT4G19230	-3.50	-3.28	-2.12	-2.49	-1.93	-2.04	-1.58	-2.10
A_84_P784432	CYP707A1	AT4G19230	-4.80	-3.82	-3.27	-3.68	-2.72	-2.72	-2.41	-3.11
A_84_P16316	CYP96A1	AT2G23180	1.97	1.91	2.68	2.61	3.28	3.41	3.82	3.79
A_84_P753234	DR356110	AT1G12672	-4.74	-4.99	-3.13	-3.74	-3.10	-3.53	-2.12	-2.76
A_84_P530411	DTA4	AT1G79760	-3.70	-3.66	-2.91	-2.35	-1.95	-1.86	-1.03	-1.43
A_84_P137739	EDL3	AT3G63060	-1.84	-2.36	-1.21	-1.36	-1.17	-1.47	-1.10	-1.20

A_84_P786422	EDL3	AT3G63060	-2.12	-2.46	-1.63	-1.80	-1.64	-1.73	-1.31	-1.58
A_84_P750050	EF182835	AT1G12670	-4.04	-4.91	-1.23	-1.70	-1.39	-1.53	0.10	-0.43
A_84_P23436	FAR1	AT5G22500	2.78	2.80	3.55	3.75	4.18	4.58	4.45	4.50
A_84_P815985	FAR1	AT5G22500	0.19	0.01	0.98	1.14	1.53	1.88	1.71	1.84
A_84_P21248	FAR8	AT3G44560	-4.67	-3.73	-3.29	-3.07	-2.52	-2.27	-2.21	-2.43
A_84_P19451	GA1	AT4G02780	-2.61	-2.21	-1.41	-1.04	-0.72	-0.34	0.95	0.68
A_84_P10329	GCL1	AT5G65280	1.76	1.78	2.85	3.48	2.66	2.89	3.87	3.76
A_84_P96816	GDU2	AT4G25760	-0.08	0.13	0.77	1.32	1.02	0.87	1.53	1.44
A_84_P15343	GH3.1	AT2G14960	-3.74	-3.18	-2.59	-2.02	1.53	1.26	2.34	1.74
A_84_P20957	GH9C2	AT1G64390	2.70	3.10	3.47	4.16	3.36	3.91	3.87	4.28
A_84_P811148	GH9C2	AT1G64390	2.15	2.63	2.92	3.59	2.85	3.33	3.25	3.69
A_84_P849563	GH9C2	AT1G64390	0.88	1.37	1.79	2.43	1.63	2.12	2.11	2.54
A_84_P811171	GH9C2	AT1G64390	-0.56	-0.40	-0.26	0.07	-0.28	-0.29	-0.06	-0.04
A_84_P802645	GH9C2	AT1G64390	-1.28	-0.61	-0.27	0.35	-0.52	0.09	0.04	0.38
A_84_P811159	GH9C2	AT1G64390	-2.20	-1.78	-1.40	-1.07	-1.73	-1.56	-1.21	-1.20
A_84_P811165	GH9C2	AT1G64390	-1.39	-0.87	-0.29	0.27	-0.70	-0.03	-0.12	0.29
A_84_P811154	GH9C2	AT1G64390	-2.05	-1.49	-0.80	-0.28	-1.22	-0.62	-0.67	-0.21
A_84_P16114	GoS3	AT1G09350	-4.98	-4.35	-3.05	-3.34	-4.16	-3.58	-2.05	-2.15
A_84_P23915	GRP2B	AT2G21060	3.49	3.87	4.28	4.73	4.26	4.60	4.90	5.33
A_84_P11248	HAI1	AT5G59220	-1.27	-1.11	0.51	0.65	0.90	0.65	0.85	0.59
A_84_P526863	IAA32	AT2G01200	-2.30	-1.63	-0.98	0.14	1.41	2.13	2.65	2.66
A_84_P24106	iqd15	AT3G49380	-3.49	-2.64	-2.56	-2.05	-2.43	-1.93	-1.80	-1.12
A_84_P823166	LAX2	AT2G21050	4.52	4.85	5.16	5.83	6.72	6.83	7.34	7.31
A_84_P11622	LAX2	AT2G21050	3.27	3.66	3.94	4.71	5.56	5.68	6.24	6.24
A_84_P55930	LFY	AT5G61850	-4.68	-3.84	-3.75	-2.50	-3.20	-2.95	-2.89	-2.46
A_84_P231259	LRP1	AT5G12330	0.71	0.81	1.43	1.88	2.12	2.22	2.17	2.39
A_84_P12315	MGL	AT1G64660	1.99	1.48	3.00	2.84	3.65	3.58	3.82	3.50
A_84_P813210	MGL	AT1G64660	-3.78	-3.92	-2.98	-2.97	-2.46	-2.10	-2.27	-2.42
A_84_P21913	MP	AT1G19850	4.15	4.38	5.57	6.11	5.24	5.61	6.25	6.59
A_84_P18699	MYB56	AT5G17800	-2.44	-2.03	-1.47	-0.68	-1.34	-0.62	-0.86	-0.40
A_84_P17140	MYC2	AT1G32640	2.99	3.11	4.28	4.31	3.82	3.99	4.20	4.05
A_84_P796568	MYC2	AT1G32640	-0.63	-0.76	-0.10	-0.71	0.07	0.20	0.72	0.40
A_84_P264370	NAC084	AT5G14000	0.65	0.65	1.39	1.25	1.40	1.50	1.88	1.67
A_84_P16775	NF-YA10	AT5G06510	-3.01	-2.96	-2.00	-1.66	-1.84	-2.03	-1.93	-1.96
A_84_P21605	NIG1	AT5G46830	-2.66	-2.58	-0.43	0.20	-1.12	-0.91	-0.92	-1.43
A_84_P19887	NIP6;1	AT1G80760	2.19	2.38	3.51	3.55	2.84	3.10	3.28	3.47
A_84_P843330	NP335312		-2.49	-2.30	-1.37	-1.64	-1.10	-1.19	-0.18	-0.82
A_84_P21121	NP454692		-1.81	-1.08	-0.47	0.56	1.91	2.53	3.02	3.02
A_84_P15935	NSP5	AT5G48180	6.70	6.26	7.05	6.75	7.72	7.87	7.81	7.75
A_84_P796347	NSP5	AT5G48180	2.31	1.85	2.96	2.53	3.46	3.60	3.65	3.55
A_84_P21273	OBP1	AT3G50410	0.65	0.87	1.10	1.82	1.76	1.87	2.02	2.18
A_84_P855408	OBP1	AT3G50410	-0.07	0.19	0.54	1.04	0.97	1.17	1.40	1.39
A_84_P11587	P5CS1	AT2G39800	0.58	0.78	1.56	1.76	1.43	1.80	1.57	1.46
A_84_P724318	PGIP1	AT5G06865	-0.49	-1.15	0.21	-0.16	0.08	-0.48	0.70	0.19
A_84_P22452	PGIP1	AT5G06860	4.25	2.86	5.16	3.48	5.57	4.45	5.73	4.61
A_84_P23398	PGIP2	AT5G06870	2.91	2.63	3.87	3.87	3.71	3.71	4.57	4.30
A_84_P787935	PIL1	AT2G46970	-2.79	-2.91	-0.91	-1.01	-1.41	-1.43	-1.21	-1.74
A_84_P521683	PIL1	AT2G46970	-3.12	-3.32	-1.35	-1.40	-1.80	-1.75	-1.61	-1.98
A_84_P111175	PIN1	AT1G73590	2.70	3.18	3.80	4.61	4.27	4.54	4.94	5.25
A_84_P13809	PRR2	AT4G13660	-2.60	-2.55	-1.08	-0.87	-1.21	-1.12	-0.74	-0.47
A_84_P178324	PUB29	AT3G18710	2.15	2.23	3.21	3.19	3.14	3.26	3.96	3.92
A_84_P811911	RAB18	AT5G66400	1.92	4.48	2.18	4.56	2.32	4.27	2.65	4.43
A_84_P861274	RAB18	AT5G66400	-1.62	-1.62	-0.03	-0.43	0.09	0.46	0.09	-0.37
A_84_P18843	RAB18	AT5G66400	-2.77	-2.43	-0.94	-1.35	-0.89	-0.63	-1.08	-1.38
A_84_P13601	RABC2b	AT3G09910	-1.64	-1.44	0.21	0.18	-0.87	-0.42	-0.08	0.02
A_84_P594466	RALFL18	AT2G33130	-3.70	-4.63	0.10	-1.85	1.23	0.40	2.99	2.47
A_84_P10392	RIC7	AT4G28560	-1.42	-0.89	-0.18	0.45	-0.73	-0.30	-0.36	0.17
A_84_P248025	RLP29	AT2G42800	0.61	1.14	1.31	2.05	2.29	2.68	2.81	3.13
A_84_P15524	RLP38	AT3G23120	-3.58	-3.57	-0.19	-1.18	-2.67	-2.96	-0.04	-1.57
A_84_P14084	ROPGEF2	AT1G01700	-2.10	-1.71	-1.27	-0.97	-1.44	-1.07	-0.67	-0.28
A_84_P753024	RTFL21	AT1G67265	-2.94	-2.78	-1.15	-1.24	-1.75	-1.27	0.09	0.08
A_84_P13455	SIP4	AT2G30360	0.73	0.78	1.53	1.44	1.86	1.75	1.95	1.92
A_84_P800880	ST2A	AT5G07010	-2.77	-3.13	-2.13	-2.60	-1.31	-0.60	-1.19	-1.07
A_84_P16777	ST2A	AT5G07010	-3.53	-3.95	-2.65	-3.20	-1.96	-1.14	-1.63	-1.48
A_84_P127161	SULTR3;1	AT3G51895	0.68	-0.01	1.55	0.48	1.95	1.61	2.22	0.97
A_84_P825382	SULTR3;1	AT3G51895	0.51	-0.26	1.53	0.41	1.82	1.35	2.12	0.79
A_84_P822209	SUS3	AT4G02280	-1.75	-2.07	-0.91	-1.09	-0.25	-0.44	-0.60	-0.85
A_84_P16611	SUS3	AT4G02280	-1.94	-2.54	-1.34	-1.44	-0.72	-0.90	-0.91	-1.19
A_84_P20094	SYPI25	AT1G11250	-4.62	-4.83	-3.31	-3.75	-3.52	-2.54	-1.92	-1.70
A_84_P821436	TA36695_3702		-4.06	-3.98	-3.47	-2.66	-3.37	-3.26	-3.27	-2.98
A_84_P825910	TA41735_3702		0.53	0.69	2.19	2.23	1.29	1.62	1.85	2.09
A_84_P833862	TA50968_3702		4.43	4.30	5.55	5.86	5.03	5.21	6.07	6.08
A_84_P836806	TA52627_3702		-4.06	-3.84	-2.76	-3.10	-2.52	-2.23	-2.15	-1.80
A_84_P768839	TC405490	AT5G66455	-0.40	-0.01	0.47	0.99	0.25	0.72	0.80	1.33

A_84_P17073	WAT1	AT1G75500	4.06	4.35	4.50	5.22	5.39	5.81	5.86	6.16
A_84_P853789	WAT1	AT1G75500	0.61	1.06	1.30	1.98	2.02	2.50	2.62	2.91
A_84_P700154	WAT1	AT1G75500	-2.77	-2.31	-2.18	-1.75	-1.60	-1.46	-1.33	-1.20
A_84_P17047	ZFP1	AT1G80730	-3.88	-2.62	-2.07	-0.90	-1.52	-1.07	-1.11	-0.54

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**Table S3. List of log<sub>2</sub> expression level of DEX-up-IAA-down genes (>1.5-fold)**

ProbeName	GeneName	AGI code	EtOH_1	EtOH_2	DEX+EtOH_1	DEX+EtOH_2	EtOH+IAA_1	EtOH+IAA_2	DEX+IAA_1	DEX+IAA_2
A_84_P12679	4CL2	AT3G21240	-0.99	-0.88	-0.12	0.11	-1.94	-1.72	-1.13	-1.17
A_84_P786411	AF418241	AT4G03060	0.49	0.97	2.35	2.63	-0.45	-0.38	1.13	1.11
A_84_P13778	AF418241	AT4G03060	-0.10	0.53	1.97	2.24	-0.78	-0.83	0.66	0.66
A_84_P23725	AHA10	AT1G17260	-2.96	-3.27	-2.33	-2.33	-3.67	-4.03	-2.78	-2.42
A_84_P159845	AT14A	AT3G28290	1.22	2.13	4.08	4.82	-0.11	0.71	3.16	3.62
A_84_P15207	AT1G14120	AT1G14120	0.40	0.72	1.20	1.61	-0.97	-0.98	-0.53	0.02
A_84_P590665	AT1G22220	AT1G22220	-1.95	-2.33	-1.00	-1.01	-3.47	-3.09	-1.92	-1.73
A_84_P20933	AT1G28610	AT1G28610	2.48	2.48	3.50	3.40	1.87	1.54	3.03	2.46
A_84_P14372	AT1G31710	AT1G31710	2.74	2.89	3.45	3.59	1.59	1.59	2.58	2.80
A_84_P10418	AT1G51440	AT1G51440	-0.37	-0.31	0.26	0.44	-1.01	-0.93	0.23	0.20
A_84_P807456	AT1G54000	AT1G54000	4.16	4.43	5.03	5.62	3.53	3.98	4.69	5.04
A_84_P223359	AT1G54000	AT1G54000	0.74	0.59	1.59	1.49	-0.29	-1.28	0.98	0.12
A_84_P807479	AT1G54000	AT1G54000	1.26	1.56	2.12	2.67	0.65	1.16	1.63	2.10
A_84_P14146	AT1G56680	AT1G56680	-3.13	-1.51	-1.37	-0.59	-4.73	-4.33	-4.45	-2.31
A_84_P753779	AT1G61795	AT1G61795	-0.06	-0.12	1.09	0.72	-1.05	-0.87	0.51	-0.01
A_84_P838837	AT1G61795	AT1G61795	-0.45	-0.50	0.64	0.24	-1.43	-1.19	0.05	-0.37
A_84_P23381	AT1G62660	AT1G62660	2.93	2.85	3.65	3.39	2.01	1.76	2.91	2.60
A_84_P804822	AT1G62660	AT1G62660	2.22	2.08	2.91	2.73	1.35	0.98	2.29	1.90
A_84_P12310	AT1G76960	AT1G76960	-1.18	-0.82	0.61	0.81	-2.13	-2.05	-1.12	-0.40
A_84_P786557	AT2G25510	AT2G25510	3.31	2.81	4.07	4.41	2.46	2.06	3.72	3.64
A_84_P60560	AT2G25510	AT2G25510	2.06	1.45	2.81	3.14	1.13	0.69	2.44	2.27
A_84_P584694	AT2G40390	AT2G40390	-2.34	-2.40	-1.23	-1.67	-3.52	-3.48	-2.11	-2.46
A_84_P22673	AT2G40435	AT2G40435	-0.65	0.41	1.35	2.61	-2.38	-0.37	-1.07	-0.42
A_84_P19250	AT3G05400	AT3G05400	0.08	0.32	0.76	1.16	-0.93	-0.50	0.76	1.36
A_84_P113022	AT3G05980	AT3G05980	0.50	1.39	1.53	2.46	-0.40	0.73	0.52	1.57
A_84_P14561	AT3G15720	AT3G15720	-0.20	0.28	0.70	0.87	-2.02	-1.27	-0.48	-0.03
A_84_P12665	AT3G16560	AT3G16560	0.79	1.13	1.47	1.99	-0.11	0.46	0.51	0.89
A_84_P832673	AT3G20340	AT3G20340	-1.13	-1.45	0.01	-0.73	-2.35	-3.08	-0.80	-1.94
A_84_P529724	AT3G20340	AT3G20340	-2.52	-2.79	-1.30	-2.04	-4.11	-3.77	-1.98	-3.41
A_84_P21186	AT3G21670	AT3G21670	0.74	0.21	2.29	1.55	-0.76	-1.67	1.09	-0.48
A_84_P18454	AT3G54600	AT3G54600	4.60	4.86	5.66	5.89	3.53	3.61	4.62	4.65
A_84_P859998	AT3G54600	AT3G54600	0.74	0.97	1.98	2.22	-0.32	-0.23	0.94	0.96
A_84_P760599	AT3G55290	AT3G55290	2.60	2.51	3.79	4.11	1.51	1.72	3.03	3.18
A_84_P108872	AT3G55290	AT3G55290	-4.59	-4.33	-2.81	-2.52	-4.94	-4.75	-3.65	-3.73
A_84_P15624	AT3G56220	AT3G56220	-0.48	0.44	0.84	2.01	-2.75	-1.04	-1.16	-0.25
A_84_P11858	AT3G59710	AT3G59710	-1.75	-1.92	0.05	0.35	-2.80	-2.51	-0.56	-0.36
A_84_P173441	AT3G61920	AT3G61920	-1.12	0.16	1.87	2.80	-2.46	-1.26	0.41	1.00
A_84_P64634	AT4G10910	AT4G10910	-0.83	-0.48	0.20	0.27	-3.95	-3.40	-2.66	-3.26
A_84_P11909	AT4G11210	AT4G11210	-0.42	-0.09	0.49	0.77	-1.86	-1.45	-0.96	-0.99
A_84_P851624	AT4G11320	AT4G11320	2.89	4.24	5.22	5.84	1.08	2.15	3.60	4.24
A_84_P20420	AT4G11320	AT4G11320	-0.14	1.26	2.42	2.96	-2.11	-0.82	0.70	1.32
A_84_P133775	AT4G16563	AT4G16563	-0.02	-0.08	1.62	0.94	-0.62	-1.44	0.54	-0.30
A_84_P12890	AT4G24050	AT4G24050	0.57	0.57	1.87	1.85	-0.82	-0.81	0.63	0.78
A_84_P15849	AT5G11930	AT5G11930	-1.21	-0.74	0.33	0.87	-3.02	-2.96	-1.85	-1.53
A_84_P13968	AT5G16410	AT5G16410	-2.29	-2.93	-1.34	-2.10	-3.62	-3.56	-2.34	-3.09
A_84_P12086	AT5G19100	AT5G19100	-1.90	-1.60	-0.95	-0.19	-2.76	-2.21	-1.59	-1.29
A_84_P831635	AT5G19100	AT5G19100	-4.13	-3.53	-3.40	-2.98	-4.65	-4.90	-3.53	-3.24
A_84_P14951	AT5G38020	AT5G38020	-1.66	-0.72	0.65	1.26	-2.80	-3.13	-1.35	-1.07
A_84_P768857	AT5G44572	AT5G44572	1.63	1.67	2.66	2.50	0.22	0.46	1.29	1.64
A_84_P528334	AT5G50915	AT5G50915	1.17	1.56	2.07	2.21	0.32	0.92	1.51	1.43
A_84_P825374	AT5G50915	AT5G50915	1.00	1.37	1.97	1.98	0.03	0.72	1.31	1.24
A_84_P784157	AT5G50915	AT5G50915	0.24	0.61	1.32	1.31	-0.53	-0.09	0.65	0.53
A_84_P15040	AT5G61660	AT5G61660	3.34	3.64	4.22	4.47	2.67	3.02	3.80	4.20
A_84_P18081	ATL8	AT1G76410	1.75	1.55	3.07	2.97	0.07	0.42	1.40	1.92
A_84_P308073	BAT5	AT4G12030	2.69	2.82	3.76	3.84	1.94	1.82	2.91	2.73
A_84_P824563	BAT5	AT4G12030	1.80	1.98	2.91	3.16	0.99	0.90	1.98	1.84
A_84_P846050	BAT5	AT4G12030	0.99	1.07	2.05	2.28	0.13	0.01	1.15	0.95
A_84_P17435	BCAT4	AT3G19710	2.48	2.84	4.53	4.73	1.56	1.49	3.49	3.87
A_84_P20230	BGLU40	AT1G26560	2.70	2.59	3.38	3.49	1.88	1.84	2.88	2.64
A_84_P22576	CUC2	AT5G53950	-0.44	0.15	0.86	1.26	-1.90	-1.10	-0.72	0.14
A_84_P23757	CUC3	AT1G76420	0.53	0.86	2.13	2.73	-0.81	-0.54	-0.23	0.58
A_84_P20172	CYP710A2	AT2G34490	2.86	3.15	3.52	3.93	1.40	1.97	2.59	2.71
A_84_P15293	CYP78A5	AT1G13710	3.68	4.28	4.42	4.90	2.81	3.42	3.19	4.22
A_84_P90469	CYP79F1	AT1G16410	1.88	2.28	4.01	4.28	1.15	1.22	3.28	3.59
A_84_P752590	CYP79F1	AT1G16410	0.90	1.40	3.21	3.38	0.17	0.25	2.26	2.54
A_84_P19484	CYP83A1	AT4G13770	5.22	5.31	6.27	6.37	4.43	4.32	5.62	5.50
A_84_P17806	DFR	AT5G42800	4.36	4.83	5.73	5.77	3.68	4.17	5.28	4.93
A_84_P845171	DFR	AT5G42800	3.69	3.90	4.95	4.92	2.95	3.50	4.56	4.17
A_84_P16271	DR4	AT1G73330	-1.53	-0.81	2.55	2.52	-3.33	-3.00	1.38	1.54
A_84_P856876	DR4	AT1G73330	-2.32	-1.86	1.70	1.47	-4.16	-4.17	0.45	0.64
A_84_P11452	ENODL6	AT1G48940	-3.52	-3.02	-2.34	-2.11	-4.56	-3.72	-2.35	-2.00
A_84_P22126	FAO3	AT3G23410	3.51	3.14	4.26	4.33	2.45	2.32	3.61	3.17

A_84_P21842	FMO GS-OX1	AT1G65860	0.75	1.08	1.69	2.07	0.15	0.09	0.77	1.10
A_84_P17028	FMO GS-OX3	AT1G62560	1.67	1.96	3.05	3.20	0.69	0.40	1.60	1.62
A_84_P760097	GRF4	AT3G52910	2.83	3.24	3.43	4.02	1.91	2.48	2.39	3.04
A_84_P20210	GSTF11	AT3G03190	1.70	2.35	2.59	2.98	0.90	0.58	1.39	1.48
A_84_P23070	HMT3	AT3G22740	-1.20	-0.55	2.71	3.14	-1.94	-3.03	1.10	1.33
A_84_P813440	IMD1	AT5G14200	7.04	7.11	7.48	7.57	6.69	6.75	6.99	7.24
A_84_P237883	IMD1	AT5G14200	4.35	4.46	5.54	5.54	3.67	3.47	4.70	4.85
A_84_P22256	IPM11	AT3G58990	2.03	2.22	3.31	3.79	1.13	1.07	2.23	2.29
A_84_P14423	IPM12	AT2G43100	4.67	4.81	5.97	6.08	4.01	3.87	5.25	5.51
A_84_P10711	JAL23	AT2G39330	0.43	-0.69	3.56	3.60	-0.91	-1.28	2.16	1.60
A_84_P22784	KNATM	AT1G14760	-1.30	-1.04	-0.48	-0.22	-1.94	-1.85	-1.03	-0.80
A_84_P861444	KNATM	AT1G14760	-0.55	-0.21	0.39	0.55	-1.45	-1.07	-0.57	-0.05
A_84_P15870	MAM1	AT5G23010	4.54	4.78	6.07	6.21	3.28	3.19	5.20	5.35
A_84_P784686	MAM1	AT5G23010	4.02	4.20	5.72	5.77	2.88	2.66	4.73	4.97
A_84_P813383	MAM1	AT5G23010	3.81	3.98	5.50	5.56	2.55	2.55	4.47	4.74
A_84_P22219	MES10	AT3G50440	4.59	4.77	6.18	6.22	3.15	3.18	4.71	4.59
A_84_P787846	MES10	AT3G50440	3.76	3.87	5.42	5.41	2.27	2.26	3.83	3.77
A_84_P13475	MGDC	AT2G11810	-2.45	-1.97	-1.31	-1.34	-3.95	-4.28	-2.19	-2.49
A_84_P189624	MLP43	AT1G70890	3.72	4.47	5.23	5.74	3.09	3.58	4.70	5.31
A_84_P18673	MYB29	AT5G07690	0.95	1.83	2.24	2.69	-1.01	-0.79	0.21	0.49
A_84_P14086	MYB59	AT5G59780	-0.30	-1.22	0.32	0.12	-1.39	-3.69	-2.01	-2.50
A_84_P14184	NAC1	AT1G56010	-1.00	-1.23	-0.40	-0.46	-3.51	-3.37	-2.85	-2.85
A_84_P13815	NLM1	AT4G19030	-2.95	-1.41	0.14	1.31	-4.02	-2.25	-0.59	0.39
A_84_P11475	PAP3	AT1G14700	3.60	3.70	4.38	4.36	3.00	3.17	4.01	3.83
A_84_P786909	PAP3	AT1G14700	3.18	3.37	4.07	3.96	2.46	2.71	3.63	3.44
A_84_P20170	PAP8	AT2G01890	1.84	1.78	4.54	3.83	-0.01	0.40	3.44	2.52
A_84_P823859	PAP8	AT2G01890	0.59	0.59	3.27	2.54	-1.06	-0.84	2.17	1.26
A_84_P11091	PLL2	AT5G02400	-3.01	-2.87	-2.13	-2.18	-3.92	-4.32	-3.47	-3.15
A_84_P23050	PR4	AT3G04720	2.86	2.40	3.77	4.07	1.49	0.34	2.62	2.44
A_84_P10135	ROT3	AT4G36380	-1.04	-0.68	0.05	0.36	-2.14	-1.67	-1.10	-1.10
A_84_P717671	RTFL6	AT4G35783	-1.68	-2.29	-1.10	-1.29	-3.45	-3.70	-2.90	-2.97
A_84_P764921	RTFL6	AT4G35783	-1.83	-2.32	-1.16	-1.33	-3.63	-3.79	-2.70	-3.26
A_84_P284250	SFP2	AT5G27360	-1.03	-0.95	0.37	0.39	-2.20	-2.14	-0.79	-0.60
A_84_P19052	TAA1	AT1G70560	4.07	4.00	4.73	5.09	2.57	2.73	3.88	4.31
A_84_P751365	TLL1	AT1G45201	4.59	4.83	5.75	6.14	3.37	3.96	4.91	4.85
A_84_P507242	TLL1	AT1G45201	2.69	2.90	3.79	4.18	1.52	2.03	2.98	2.94
A_84_P116602	WOL	AT2G01830	2.01	2.42	2.95	3.20	1.04	1.52	1.69	1.91
A_84_P784848	WOL	AT2G01830	-0.21	0.42	0.95	1.27	-0.84	-0.58	-0.25	-0.04
A_84_P22412	WRKY13	AT4G39410	-1.76	-1.06	-0.85	-0.29	-2.52	-1.92	-1.32	-0.91
A_84_P544339	XND1	AT5G64530	-0.83	-1.07	0.55	0.28	-1.65	-1.95	-0.18	-0.69
A_84_P18684	YUC4	AT5G11320	-0.16	0.54	0.54	1.37	-1.65	-1.54	-1.99	-0.35
A_84_P767924	YUC4	AT5G11320	-0.61	-0.10	-0.04	0.88	-2.39	-2.17	-2.09	-0.92
A_84_P591116	ZPR2	AT3G60890	-1.24	-0.98	0.11	0.42	-2.22	-1.99	-0.99	-0.23
A_84_P13717	ZPR3	AT3G52770	-1.28	-1.11	-0.07	0.10	-2.12	-1.77	-0.57	0.21

**Table S4. List of log<sub>2</sub> expression level of DEX-down-IAA-up genes (>1.5-fold)**

ProbeName	GeneName	AGI code	EtOH_1	EtOH_2	DEX+EtOH_1	DEX+EtOH_2	EtOH+IAA_1	EtOH+IAA_2	DEX+IAA_1	DEX+IAA_2
A_84_P52180	AGL8	AT5G60910	-2.43	-2.01	-3.81	-2.61	-1.05	-0.92	-1.51	-1.36
A_84_P821066	AGL8	AT5G60910	-2.68	-2.19	-4.36	-2.83	-1.09	-1.26	-2.02	-1.87
A_84_P208088	AGP4	AT5G10430	4.03	3.99	2.89	2.64	4.73	4.86	3.73	3.78
A_84_P851051	AGP4	AT5G10430	0.97	1.00	-0.27	-0.26	1.73	1.96	0.75	0.96
A_84_P524502	ARGOS	AT3G59900	4.31	3.98	2.50	1.39	5.30	5.08	4.35	4.39
A_84_P18985	AT1G11740	AT1G11740	-1.30	-1.00	-2.37	-2.03	0.39	0.69	-0.63	-0.53
A_84_P751955	AT1G12080	AT1G12080	4.53	5.29	2.72	3.37	5.62	5.95	4.48	4.86
A_84_P848382	AT1G12080	AT1G12080	4.26	4.81	2.48	3.13	5.34	5.64	4.19	4.58
A_84_P807614	AT1G12080	AT1G12080	2.41	2.97	0.49	1.30	3.42	3.81	2.27	2.71
A_84_P603116	AT1G13670	AT1G13670	-0.36	-0.27	-2.70	-2.52	1.37	1.29	0.41	1.02
A_84_P22857	AT1G18020	AT1G17990	3.98	3.07	2.10	0.95	5.08	4.04	3.09	2.22
A_84_P803569	AT1G18020	AT1G18020	3.15	2.21	1.32	0.08	4.19	3.19	2.29	1.39
A_84_P853555	AT1G18020	AT1G18020	1.53	0.40	-0.29	-1.57	2.45	1.52	0.55	-0.44
A_84_P23662	AT1G23040	AT1G23040	2.49	2.51	1.66	1.65	3.28	3.30	2.50	2.59
A_84_P858244	AT1G23040	AT1G23040	-1.10	-1.30	-1.87	-1.75	-0.22	-0.25	-0.97	-0.87
A_84_P279980	AT1G29440	AT1G29440	2.24	2.37	-1.90	-1.75	3.68	3.48	2.19	2.26
A_84_P19713	AT1G29450	AT1G29450	1.12	1.36	-2.44	-2.20	2.78	2.66	1.84	1.79
A_84_P15930	AT1G29460	AT1G29460	-0.23	0.14	-4.86	-3.65	1.92	1.98	1.09	1.34
A_84_P754333	AT1G29465	AT1G29465	4.46	4.25	3.55	3.31	5.14	5.01	4.78	4.56
A_84_P843096	AT1G29490	AT1G29490	-2.42	-2.09	-3.63	-4.72	1.40	1.31	0.62	0.75
A_84_P11207	AT1G29500	AT1G29500	1.56	1.55	-0.20	0.18	3.14	3.00	2.43	2.39
A_84_P16068	AT1G33790	AT1G33790	2.86	2.87	1.22	0.83	3.88	4.04	2.26	2.27
A_84_P13577	AT1G36940	AT1G36940	2.17	1.86	1.42	1.15	3.13	3.15	2.65	2.76
A_84_P19990	AT1G68810	AT1G68810	0.78	0.83	-0.90	-0.32	1.81	1.68	0.68	0.49
A_84_P19196	AT1G75490	AT1G75490	1.05	-0.31	0.11	-1.31	1.84	1.14	1.80	1.18
A_84_P795438	AT1G75490	AT1G75490	-3.52	-4.98	-2.52	-4.94	-2.57	-3.10	-2.90	-3.48
A_84_P118432	AT1G78100	AT1G78100	2.66	2.57	1.72	1.70	3.60	3.71	3.04	3.09
A_84_P758488	AT2G18969	AT2G18969	1.11	1.36	-0.40	-0.18	2.74	2.89	1.41	1.63
A_84_P595309	AT2G18970	AT2G18970	-3.14	-2.68	-4.40	-4.66	-1.53	-1.29	-2.96	-2.67
A_84_P136225	AT2G32560	AT2G32560	2.70	2.72	0.29	0.13	3.44	3.70	2.57	2.57
A_84_P835471	AT2G34080	AT2G34080	0.00	-0.43	-1.29	-2.03	0.72	0.77	-0.25	-0.57
A_84_P17235	AT2G34080	AT2G34080	-0.72	-1.18	-1.98	-3.25	-0.04	-0.02	-1.01	-1.28
A_84_P21046	AT2G38600	AT2G38600	-3.96	-3.40	-4.62	-4.97	-2.19	-2.21	-3.67	-3.76
A_84_P13439	AT2G39980	AT2G39980	5.22	4.28	3.59	2.55	6.52	6.29	5.32	5.12
A_84_P196604	AT2G40530	AT2G40530	-1.23	-1.04	-2.36	-2.03	-0.20	0.14	-1.04	-0.84
A_84_P173611	AT2G41170	AT2G41170	0.81	0.71	0.06	-0.13	2.07	1.89	1.36	1.25
A_84_P297904	AT2G41230	AT2G41230	4.84	3.53	4.05	2.17	5.78	5.48	5.85	5.78
A_84_P795880	AT2G41231	AT2G41231	0.16	-1.07	-0.68	-2.28	0.95	0.80	1.13	1.14
A_84_P160753	AT2G41800	AT2G41800	-2.08	-0.34	-3.53	-2.17	0.59	1.52	-0.66	0.02
A_84_P757973	AT2G44578	AT2G44578	-1.82	-2.39	-2.52	-3.52	-0.09	-0.98	-0.94	-1.60
A_84_P21135	AT3G03820	AT3G03820	0.97	1.34	0.12	0.11	2.49	2.38	1.75	1.85
A_84_P20189	AT3G03830	AT3G03830	-0.86	-0.56	-2.18	-2.02	1.41	1.37	0.69	0.87
A_84_P18297	AT3G03840	AT3G03840	-2.85	-2.45	-3.64	-3.29	-0.96	-1.09	-1.60	-1.15
A_84_P554482	AT3G03850	AT3G03850	-0.67	-0.31	-1.95	-1.71	1.12	1.10	0.57	0.87
A_84_P20213	AT3G04000	AT3G04000	3.47	1.70	2.05	0.89	4.13	2.29	2.45	0.93
A_84_P793541	AT3G04000	AT3G04000	1.20	-0.67	-0.17	-1.21	1.95	0.07	0.35	-1.19
A_84_P18298	AT3G05150	AT3G05150	-1.87	-1.86	-3.40	-2.55	-0.79	-0.51	-0.93	-0.76
A_84_P832301	AT3G05150	AT3G05150	-2.97	-2.55	-3.37	-3.33	-1.47	-1.25	-1.74	-1.36
A_84_P14558	AT3G22820	AT3G22820	0.29	0.22	-1.48	-1.69	1.27	1.06	0.14	0.01
A_84_P21181	AT3G26760	AT3G26760	0.26	0.75	-1.83	-0.99	1.78	2.16	0.89	1.32
A_84_P558998	AT3G42800	AT3G42800	-0.69	-0.22	-1.92	-1.21	1.07	1.19	0.86	1.17
A_84_P17461	AT3G43430	AT3G43430	1.15	-1.17	-0.33	-0.93	1.84	1.99	0.70	0.84
A_84_P860746	AT3G47510	AT3G47510	0.90	0.80	-0.69	-0.77	2.18	2.26	1.22	1.08
A_84_P556364	AT3G47510	AT3G47510	-1.03	-1.02	-2.59	-2.90	0.37	0.35	-0.73	-0.64
A_84_P509388	AT3G52520	AT3G52520	-3.31	-3.70	-4.04	-4.40	-2.14	-2.86	-2.87	-4.48
A_84_P13751	AT3G60490	AT3G60490	-0.13	-0.22	-1.28	-1.49	0.47	0.54	-0.52	-0.39
A_84_P20398	AT4G03140	AT4G03140	1.21	0.97	-1.15	-0.70	3.27	3.26	1.46	1.79
A_84_P848192	AT4G03140	AT4G03140	-2.04	-1.99	-4.06	-4.15	0.31	0.42	-1.47	-1.17
A_84_P19468	AT4G09890	AT4G09890	2.68	2.91	1.37	1.79	3.65	3.80	3.06	3.10
A_84_P764623	AT4G15248	AT4G15248	-0.37	-1.64	-1.04	-2.44	1.62	0.19	0.47	-0.80
A_84_P572494	AT4G28703	AT4G28703	-0.31	-0.33	-1.36	-1.89	1.50	1.06	-0.32	-0.27
A_84_P787406	AT4G28703	AT4G28703	-0.84	-0.78	-1.97	-2.30	0.88	0.53	-1.07	-0.80
A_84_P831098	AT4G33800	AT4G33800	-2.72	-2.74	-3.97	-3.53	-2.11	-1.68	-2.37	-1.93
A_84_P222749	AT4G33800	AT4G33800	-2.69	-2.56	-3.62	-3.44	-1.93	-1.84	-2.38	-2.04
A_84_P21449	AT4G34770	AT4G34770	3.51	3.58	0.84	0.55	4.21	4.30	3.29	3.55
A_84_P13885	AT4G35190	AT4G35190	-0.18	-0.52	-2.35	-1.81	0.56	0.33	-0.79	-0.67
A_84_P797355	AT4G36110	AT4G36110	0.15	0.57	-1.19	-0.81	1.26	1.15	0.72	0.87
A_84_P17680	AT4G38860	AT4G38860	1.12	1.02	0.03	-0.01	1.99	1.98	1.77	1.78
A_84_P768708	AT5G01210	AT5G01215	-1.24	-0.87	-1.31	-0.53	-0.50	-0.18	-0.17	0.03
A_84_P707180	AT5G01210	AT5G01215	-2.00	-1.97	-2.45	-2.20	-1.54	-1.17	-1.59	-1.24
A_84_P17703	AT5G01210	AT5G01210	5.09	4.96	4.35	3.86	6.11	6.14	6.13	6.16
A_84_P22435	AT5G01740	AT5G01740	2.88	2.14	0.57	-0.26	4.44	4.46	2.78	2.60



A_84_P16764	AT5G02750	AT5G02750	0.22	0.26	-1.55	-2.12	1.12	1.11	-0.15	-0.21
A_84_P17710	AT5G02760	AT5G02760	4.79	4.16	2.90	2.86	6.95	7.20	6.86	6.90
A_84_P172771	AT5G06930	AT5G06930	0.95	1.09	0.11	0.19	1.84	1.82	1.46	1.59
A_84_P831040	AT5G06930	AT5G06930	-2.98	-2.76	-3.52	-3.41	-2.22	-2.06	-2.44	-2.42
A_84_P196504	AT5G18010	AT5G18010	-0.05	0.40	-2.55	-1.91	2.29	2.47	1.41	1.60
A_84_P147028	AT5G18020	AT5G18020	0.26	0.87	-1.04	-0.94	2.40	2.44	1.94	2.01
A_84_P152798	AT5G18030	AT5G18030	0.62	1.06	-0.67	-0.40	2.88	2.98	2.53	2.64
A_84_P141269	AT5G18050	AT5G18050	1.07	1.68	-0.27	-0.01	3.74	3.93	3.20	3.41
A_84_P94979	AT5G18060	AT5G18060	0.68	0.86	-0.69	-0.59	2.95	3.05	2.45	2.59
A_84_P272980	AT5G18080	AT5G18080	0.38	0.93	-0.72	-0.96	2.89	3.01	2.47	2.54
A_84_P79425	AT5G20820	AT5G20820	2.17	2.33	0.85	0.94	3.52	3.69	2.66	3.08
A_84_P824903	AT5G22460	AT5G22460	1.56	1.60	0.40	0.32	2.21	2.20	2.07	1.53
A_84_P167453	AT5G22460	AT5G22460	0.96	0.95	-0.32	-0.37	1.65	1.55	1.35	0.94
A_84_P824908	AT5G22460	AT5G22460	-2.82	-2.49	-3.43	-4.04	-1.76	-1.90	-2.06	-2.69
A_84_P13987	AT5G25190	AT5G25190	3.01	2.84	0.84	0.74	4.33	4.35	3.20	3.27
A_84_P194934	AT5G52870	AT5G52870	3.24	3.15	2.37	2.27	3.91	3.97	3.57	3.51
A_84_P214628	AT5G52900	AT5G52900	1.71	0.53	-1.35	-1.02	4.17	3.88	4.12	3.38
A_84_P507058	AT5G57760	AT5G57760	5.23	4.51	3.37	2.18	5.92	6.27	5.49	5.42
A_84_P789543	AT5G57760	AT5G57760	4.05	3.30	2.13	1.08	4.78	5.01	4.27	4.23
A_84_P23538	AT5G58620	AT5G58620	0.92	0.75	0.23	0.06	1.67	1.41	1.22	1.12
A_84_P79759	AT5G62280	AT5G62280	0.75	-0.79	-2.07	-2.03	3.43	3.16	3.53	3.24
A_84_P54620	AT5G64780	AT5G64780	0.70	0.29	-1.10	-0.93	1.37	1.28	0.46	0.60
A_84_P14971	ATAUX2-11	AT5G43700	3.59	3.58	2.73	2.82	4.55	4.78	4.32	4.45
A_84_P818769	ATAUX2-11	AT5G43700	1.60	1.61	0.73	0.79	2.55	2.83	2.30	2.46
A_84_P20866	AXR3	AT1G04250	3.82	3.80	2.93	2.79	4.63	4.70	4.28	4.33
A_84_P838052	AXR3	AT1G04250	1.47	1.55	0.61	0.52	2.26	2.34	1.84	1.99
A_84_P18482	BRH1	AT3G61460	5.33	5.27	4.61	4.67	6.12	6.32	5.92	5.90
A_84_P22483	CAX7	AT5G17860	3.85	2.96	3.22	1.49	4.85	3.93	4.03	2.59
A_84_P592126	CCD8	AT4G32810	-2.62	-2.91	-4.62	-4.11	-0.72	-1.03	-1.65	-2.27
A_84_P14421	CKX1	AT2G41510	-1.81	-1.55	-3.71	-2.52	-0.27	-0.26	-1.67	-1.85
A_84_P302670	CLE22	AT5G12235	-1.61	-1.54	-2.45	-2.43	-0.84	-0.65	-1.29	-1.24
A_84_P762938	DQ108837	AT4G22545	-2.24	-2.01	-3.79	-3.66	-0.62	-0.58	-1.38	-1.53
A_84_P762938	DQ108837	AT4G22545	-2.04	-2.22	-4.22	-4.40	-0.53	-0.52	-1.65	-1.41
A_84_P762938	DQ108837	AT4G22545	-2.15	-1.98	-4.50	-4.13	-0.53	-0.51	-1.87	-1.60
A_84_P762938	DQ108837	AT4G22545	-2.13	-2.17	-4.41	-4.74	-0.51	-0.57	-1.79	-1.38
A_84_P762938	DQ108837	AT4G22545	-2.60	-2.12	-4.19	-4.48	-0.61	-0.80	-1.97	-1.75
A_84_P298034	ENODL18	AT1G08500	0.30	0.44	-0.35	-0.74	0.93	1.17	0.15	0.14
A_84_P23714	ESC	AT1G20900	1.71	1.55	-0.49	-0.59	3.47	3.64	2.21	2.48
A_84_P806966	EXPA8	AT2G40610	2.55	2.61	1.18	1.33	3.41	3.59	2.92	3.32
A_84_P11569	EXPA8	AT2G40610	1.80	1.87	0.55	0.74	2.70	2.96	2.27	2.67
A_84_P788919	EXPA8	AT2G40610	-2.74	-2.55	-3.94	-3.83	-1.83	-1.53	-2.01	-1.60
A_84_P16678	GA200X1	AT4G25420	-0.88	-0.57	-1.97	-2.01	1.52	1.81	1.41	1.53
A_84_P18218	HB17	AT2G01430	-2.76	-2.63	-3.66	-3.48	0.45	0.38	-0.29	-0.81
A_84_P14556	IAA19	AT3G15540	1.15	0.96	-1.18	-0.12	5.74	5.81	5.12	5.25
A_84_P820504	IAA2	AT3G23030	5.32	4.83	2.96	2.56	7.94	7.72	7.64	7.40
A_84_P19308	IAA2	AT3G23030	2.76	2.21	0.56	0.13	5.47	5.25	5.23	5.01
A_84_P55550	IAA29	AT4G32280	-1.01	-1.85	-2.92	-2.75	4.74	4.55	4.82	4.59
A_84_P13379	IAA6	AT1G52830	-1.95	-1.25	-4.33	-4.11	2.23	1.99	0.66	-0.11
A_84_P761007	IAA7	AT3G23050	5.01	4.95	4.24	4.23	6.69	6.80	6.50	6.51
A_84_P760471	IAA7	AT3G23050	-2.81	-2.78	-2.77	-3.27	-0.60	-0.79	-0.73	-0.93
A_84_P544465	KDR	AT1G26945	3.64	3.22	1.13	1.35	6.53	6.43	5.63	5.77
A_84_P789302	KDR	AT1G26945	-1.12	-1.43	-2.84	-3.10	1.96	1.99	1.07	1.27
A_84_P16293	LAC6	AT2G46570	-0.55	-0.44	-2.16	-1.16	1.52	1.36	0.74	0.17
A_84_P15136	LRX2	AT1G62440	-3.27	-3.19	-3.95	-4.26	-1.31	-2.09	-3.07	-3.13
A_84_P10579	LUP1	AT1G78970	2.74	2.67	1.34	1.16	4.30	4.20	3.54	3.49
A_84_P174971	NPY5	AT4G37590	3.59	3.60	2.50	2.79	4.42	4.38	3.88	3.80
A_84_P784468	NPY5	AT4G37590	3.68	3.72	2.65	3.03	4.45	4.44	3.98	3.92
A_84_P19825	PKS4	AT5G04190	1.83	1.18	-1.39	-2.21	2.80	2.52	1.60	1.86
A_84_P173491	RGF9	AT5G64770	2.72	2.60	0.52	0.91	4.09	3.89	2.93	2.93
A_84_P19638	ROXY2	AT5G14070	-3.71	-2.22	-4.47	-3.35	-1.96	-1.34	-1.51	-1.06
A_84_P519104	RTFL16	AT3G25717	2.36	2.74	1.16	1.72	3.54	3.61	3.46	3.51
A_84_P16734	SAUR15	AT4G38850	0.12	0.43	-3.79	-3.52	2.18	2.21	1.33	1.48
A_84_P22555	SAUR68	AT1G29510	2.07	2.15	0.33	0.85	3.03	3.02	2.24	2.43
A_84_P15234	sepB1	AT1G11080	0.21	0.43	-0.39	-1.67	1.04	1.24	-0.70	-0.22
A_84_P266190	SOFL1	AT1G26210	1.68	1.50	-0.06	0.10	2.31	2.60	1.00	0.95
A_84_P753467	SOFL1	AT1G26208	-0.69	-0.92	-2.04	-1.75	-0.09	-0.04	-1.04	-0.90
A_84_P23812	ST4C	AT1G13430	-3.18	-2.66	-3.85	-3.34	-1.36	-1.20	-1.87	-1.24
A_84_P790823	WVD2	AT5G28646	-2.81	-2.75	-4.26	-4.31	-0.47	-1.31	-2.04	-2.68
A_84_P10083	XTH19	AT4G30290	-1.51	-1.99	-3.59	-3.09	1.91	1.73	-0.17	-0.37
A_84_P15301	ZF14	AT1G58340	2.91	3.08	1.65	1.87	3.83	3.99	3.31	3.18

**Table S5. List of log<sub>2</sub> expression level of DEX-down-IAA-down genes (>1.5-fold)**

ProbeName	GeneName	AGI code	EtOH_1	EtOH_2	DEX+EtOH_1	DEX+EtOH_2	EtOH+IAA_1	EtOH+IAA_2	DEX+IAA_1	DEX+IAA_2
A_84_P22208	ABCA4	AT3G47750	-0.53	-0.50	-1.23	-1.54	-1.32	-1.12	-1.74	-1.88
A_84_P14374	ABCG14	AT1G31770	2.53	2.86	1.32	1.85	1.27	1.65	-0.07	0.82
A_84_P12588	ACL	AT2G23600	5.10	4.94	2.41	2.30	3.26	2.95	1.05	0.99
A_84_P15145	ACR3	AT1G76990	5.86	5.43	4.82	4.31	5.16	4.45	4.35	3.63
A_84_P808028	ACR3	AT1G76990	-1.80	-2.13	-2.75	-3.39	-3.20	-3.41	-3.68	-3.50
A_84_P123362	AFP2	AT1G13740	2.19	2.29	0.16	0.39	1.44	1.22	-0.55	-0.42
A_84_P298854	AHL20	AT4G14465	-0.28	-0.01	-2.08	-1.71	-2.78	-1.99	-3.54	-2.08
A_84_P753477	AK220633	AT1G68568	-0.80	-0.91	-1.77	-1.84	-2.03	-2.22	-3.06	-3.55
A_84_P23733	APT2	AT1G80050	3.48	3.80	2.10	2.43	1.46	1.93	0.70	1.00
A_84_P17147	ARAB-1	AT1G28670	3.72	3.89	2.74	3.09	2.62	3.17	2.34	2.70
A_84_P10395	ARR7	AT1G19050	3.63	3.62	2.52	1.80	1.87	1.58	1.46	0.63
A_84_P15163	AS2	AT1G65620	1.11	1.44	-0.27	0.19	0.27	0.74	-0.81	-0.18
A_84_P826615	AS2	AT1G65620	-3.03	-2.17	-3.88	-3.56	-3.28	-2.94	-4.96	-3.89
A_84_P18932	AST91	AT1G23090	2.68	2.40	-2.17	-2.15	1.80	1.60	-1.65	-1.40
A_84_P822023	AST91	AT1G23090	0.24	0.07	-4.45	-3.75	-0.74	-0.84	-3.90	-4.34
A_84_P12740	AT1G02630	AT1G02630	-2.77	-1.94	-4.74	-4.47	-3.46	-2.75	-3.71	-3.51
A_84_P21249	AT1G02650	AT1G02650	-0.94	-1.25	-2.77	-2.94	-1.71	-2.09	-3.38	-3.32
A_84_P17060	AT1G09440	AT1G09440	1.63	1.65	0.69	0.61	0.43	0.60	-0.08	-0.02
A_84_P22769	AT1G11700	AT1G11700	4.90	4.97	3.70	3.94	3.58	3.97	2.91	3.11
A_84_P752662	AT1G12064	AT1G12064	0.78	0.81	-0.43	-0.68	-0.01	-0.04	-1.29	-0.97
A_84_P21881	AT1G12100	AT1G12100	0.71	0.88	-0.22	0.26	-0.22	0.22	-0.71	-0.02
A_84_P20792	AT1G15670	AT1G15670	3.09	2.24	1.34	0.96	1.75	0.94	0.93	0.77
A_84_P854414	AT1G15670	AT1G15670	0.78	0.01	-0.81	-1.29	-0.57	-1.28	-1.30	-1.52
A_84_P516067	AT1G17345	AT1G17345	-1.67	-1.80	-2.67	-2.50	-2.87	-2.60	-3.14	-2.82
A_84_P20019	AT1G17960	AT1G17960	2.55	1.38	1.20	-0.83	1.87	0.62	1.14	-1.27
A_84_P14276	AT1G22550	AT1G22550	2.29	2.48	1.45	1.66	1.26	1.85	1.10	1.31
A_84_P13332	AT1G22570	AT1G22570	1.85	1.80	-1.93	-1.47	0.29	0.09	-1.94	-1.82
A_84_P18707	AT1G23390	AT1G23390	3.24	2.47	2.12	1.79	2.13	1.21	1.38	0.97
A_84_P786694	AT1G23390	AT1G23390	3.12	2.33	2.07	1.59	1.96	1.12	1.31	0.92
A_84_P869207	AT1G23390	AT1G23390	2.57	1.77	1.46	0.93	1.46	0.55	0.75	0.26
A_84_P796487	AT1G23390	AT1G23390	-0.39	-1.05	-1.26	-1.84	-1.64	-2.32	-2.14	-2.58
A_84_P16262	AT1G25230	AT1G25230	4.46	4.54	2.25	1.83	3.16	3.55	1.45	1.79
A_84_P785932	AT1G25400	AT1G25400	2.50	1.39	0.92	-0.82	1.59	0.21	0.66	-0.38
A_84_P67084	AT1G25400	AT1G25400	1.32	0.03	-0.29	-2.21	0.20	-1.09	-0.65	-1.66
A_84_P15298	AT1G25520	AT1G25520	2.31	1.94	0.28	0.65	1.48	1.19	0.20	0.58
A_84_P10039	AT1G27020	AT1G27020	4.97	4.57	0.54	0.19	3.49	2.66	-0.77	-1.42
A_84_P860087	AT1G27020	AT1G27020	4.30	3.86	-0.10	-0.30	2.91	1.96	-1.16	-1.64
A_84_P200014	AT1G28660	AT1G28660	3.34	3.36	2.60	2.43	2.35	2.51	2.23	2.23
A_84_P15451	AT1G44760	AT1G44760	5.01	5.39	4.42	4.63	4.22	4.69	3.84	3.99
A_84_P816367	AT1G44760	AT1G44760	4.54	4.89	3.90	4.13	3.77	4.07	3.30	3.51
A_84_P16258	AT1G49230	AT1G49230	2.05	2.55	-1.22	-1.17	1.39	1.86	-0.71	0.09
A_84_P66364	AT1G49310	AT1G49310	-0.44	-0.05	-2.51	-3.16	-1.70	-1.41	-3.71	-2.89
A_84_P22625	AT1G52270	AT1G52270	1.86	2.25	1.18	1.44	1.08	1.63	1.34	1.67
A_84_P14112	AT1G52290	AT1G52290	0.19	0.35	-1.40	-1.14	-1.14	-1.28	-2.00	-1.97
A_84_P14201	AT1G60680	AT1G60680	-2.07	-1.31	-2.88	-2.64	-3.11	-2.42	-3.42	-3.56
A_84_P184384	AT1G68440	AT1G68440	4.71	3.79	2.19	1.18	4.05	2.66	1.58	0.55
A_84_P249125	AT1G68500	AT1G68500	2.52	2.47	-0.03	0.00	1.73	1.75	0.21	0.19
A_84_P23779	AT1G68520	AT1G68520	5.03	4.20	2.01	1.77	3.32	3.29	0.30	-0.34
A_84_P18751	AT1G68570	AT1G68570	4.22	4.05	3.23	3.30	3.04	2.79	2.21	1.73
A_84_P150878	AT1G69760	AT1G69760	3.45	3.21	0.97	1.00	2.56	2.30	0.82	1.01
A_84_P556074	AT1G70985	AT1G70985	1.69	1.59	0.62	0.48	0.86	0.78	0.71	0.54
A_84_P789698	AT1G72416	AT1G72416	0.85	0.72	-2.86	-2.43	-2.26	-1.59	-2.98	-1.93
A_84_P715317	AT1G72416	AT1G72416	0.36	0.23	-4.29	-3.19	-2.50	-2.43	-4.03	-2.74
A_84_P22853	AT1G75960	AT1G75960	0.38	0.34	-1.80	-1.07	-2.23	-1.95	-2.25	-1.76
A_84_P145909	AT1G78020	AT1G78020	2.37	2.58	0.66	1.04	1.01	1.16	-0.75	-0.84
A_84_P14286	AT1G78260	AT1G78260	3.45	3.55	2.28	2.80	2.58	2.80	1.91	2.41
A_84_P751743	AT1G78260	AT1G78260	2.67	2.95	1.53	2.07	1.80	2.06	1.07	1.58
A_84_P260410	AT1G78450	AT1G78450	-0.67	-0.14	-2.37	-1.96	-4.34	-3.01	-3.97	-3.42
A_84_P12443	AT1G80440	AT1G80440	4.44	3.40	3.07	2.31	3.17	0.96	1.78	0.87
A_84_P857022	AT1G80440	AT1G80440	1.33	0.28	-0.06	-0.77	-0.12	-2.31	-1.38	-2.39
A_84_P268090	AT2G01300	AT2G01300	-0.34	-0.41	-1.22	-1.84	-2.13	-2.91	-2.09	-2.49
A_84_P505592	AT2G04795	AT2G04795	1.27	0.55	-1.72	-2.53	0.14	-0.76	-2.12	-2.35
A_84_P116392	AT2G15910	AT2G15910	-3.40	-3.56	-3.89	-3.63	-3.43	-3.67	-3.86	-4.24
A_84_P843500	AT2G15910	AT2G15910	-3.62	-3.89	-4.40	-4.56	-4.28	-4.64	-4.16	-3.24
A_84_P235483	AT2G18360	AT2G18360	0.46	0.34	-2.07	-2.24	-0.98	-0.82	-2.21	-1.83
A_84_P132645	AT2G20670	AT2G20670	5.90	4.91	4.52	3.93	3.49	2.82	2.82	2.22
A_84_P513598	AT2G22122	AT2G22122	1.14	1.25	-0.20	-0.16	0.32	0.60	0.41	0.38
A_84_P21994	AT2G26440	AT2G26440	2.92	2.97	0.28	0.06	2.31	2.35	0.54	0.42
A_84_P12501	AT2G28200	AT2G28200	0.70	0.20	-0.42	-0.79	-0.45	-0.50	-0.91	-1.13
A_84_P12493	AT2G30540	AT2G30540	-1.06	-0.46	-2.20	-2.10	-2.19	-2.08	-3.13	-2.14
A_84_P304680	AT2G30600	AT2G30600	5.00	4.97	3.10	3.12	3.87	3.44	2.79	3.03
A_84_P189284	AT2G30930	AT2G30930	2.92	2.83	2.28	1.61	2.00	1.59	1.67	1.08

A_84_P23890	AT2G32960	AT2G32960	2.55	2.22	0.26	-0.08	1.56	1.29	-0.76	-0.83
A_84_P565934	AT2G34010	AT2G34010	-1.30	-1.29	-2.69	-2.43	-3.41	-3.17	-3.51	-3.20
A_84_P11644	AT2G37460	AT2G37460	0.47	0.62	-2.41	-2.66	-1.46	-1.24	-3.77	-2.75
A_84_P150068	AT2G43140	AT2G43140	2.07	1.16	-1.53	-1.80	-1.51	-3.19	-2.53	-1.79
A_84_P838362	AT2G43140	AT2G43140	0.67	-0.13	-2.43	-2.38	-2.27	-4.08	-3.27	-2.38
A_84_P809606	AT2G45180	AT2G45180	9.21	9.01	7.09	7.05	7.88	7.94	6.33	6.37
A_84_P272390	AT2G45180	AT2G45180	8.64	8.43	6.49	6.47	7.25	7.33	5.73	5.79
A_84_P809597	AT2G45180	AT2G45180	8.40	8.14	6.16	6.10	6.96	7.13	5.41	5.52
A_84_P801096	AT2G45180	AT2G45180	8.32	8.11	6.13	6.07	6.93	7.01	5.35	5.45
A_84_P597183	AT2G45900	AT2G45900	1.40	1.27	0.45	0.64	0.01	0.41	-0.54	-0.65
A_84_P144789	AT3G01860	AT3G01860	3.38	3.30	1.84	2.10	2.39	2.49	1.86	1.86
A_84_P829153	AT3G01860	AT3G01860	2.62	2.56	1.06	1.35	1.61	1.76	1.09	1.09
A_84_P520804	AT3G06020	AT3G06020	-0.95	-1.02	-1.70	-1.97	-2.90	-2.25	-2.72	-2.83
A_84_P108832	AT3G06070	AT3G06070	1.02	1.00	-1.11	-1.26	0.20	0.15	-0.99	-1.21
A_84_P242653	AT3G06990	AT3G06990	-2.36	-1.97	-3.02	-3.28	-3.41	-3.22	-3.80	-3.16
A_84_P21242	AT3G11280	AT3G11280	3.56	3.13	2.48	2.04	2.37	2.10	1.60	1.12
A_84_P810292	AT3G11930	AT3G11930	3.75	3.15	3.24	2.86	2.93	2.92	2.44	2.16
A_84_P22107	AT3G11930	AT3G11930	2.64	2.07	2.22	1.86	1.89	1.92	1.33	1.18
A_84_P861062	AT3G11930	AT3G11930	1.99	1.58	1.63	1.38	1.41	1.24	0.79	0.54
A_84_P857510	AT3G11930	AT3G11930	-0.53	-1.64	-1.90	-2.38	-1.80	-2.33	-4.54	-4.30
A_84_P16506	AT3G14770	AT3G14770	2.00	1.33	0.84	0.59	1.07	0.61	0.14	-0.15
A_84_P823098	AT3G14770	AT3G14770	-0.17	-0.55	-1.14	-1.54	-0.87	-1.32	-1.71	-1.94
A_84_P17452	AT3G14780	AT3G14780	-0.71	-0.63	-1.47	-1.72	-1.66	-1.47	-1.79	-2.22
A_84_P296824	AT3G15450	AT3G15450	8.08	7.93	6.88	6.97	6.82	6.19	6.17	6.36
A_84_P855440	AT3G15450	AT3G15450	5.89	5.58	4.90	4.94	4.75	3.91	4.06	4.30
A_84_P805118	AT3G15450	AT3G15450	6.08	5.68	4.83	4.87	4.76	4.08	3.97	4.27
A_84_P805135	AT3G15450	AT3G15450	5.41	4.94	4.14	4.10	4.03	3.25	3.33	3.57
A_84_P818968	AT3G15630	AT3G15630	2.33	2.64	1.29	1.78	1.24	1.77	0.88	1.74
A_84_P81409	AT3G15630	AT3G15630	2.49	2.50	1.17	1.51	0.35	0.76	0.15	0.90
A_84_P582090	AT3G18773	AT3G18773	-0.12	-0.89	-2.90	-3.37	-1.30	-1.97	-3.91	-3.42
A_84_P831705	AT3G18900	AT3G18900	0.07	0.44	-1.15	-0.93	-1.58	-1.16	-3.19	-2.00
A_84_P587454	AT3G19850	AT3G19850	2.28	1.76	-0.42	-0.98	-0.38	0.07	-1.51	-1.72
A_84_P591022	AT3G21530	AT3G21530	-0.35	-0.83	-1.50	-1.90	-1.33	-1.73	-2.06	-2.67
A_84_P13632	AT3G26700	AT3G26700	1.53	1.81	0.59	1.14	0.69	0.83	0.70	0.87
A_84_P826141	AT3G27030	AT3G27030	1.97	1.96	-0.19	-0.50	0.92	1.19	-0.86	-0.74
A_84_P534191	AT3G27030	AT3G27030	0.99	0.94	-1.26	-1.35	-0.04	0.12	-1.98	-1.98
A_84_P21738	AT3G28345	AT3G28345	1.57	1.96	0.25	0.34	0.55	0.76	-0.51	-0.04
A_84_P832213	AT3G28345	AT3G28345	0.45	0.86	-0.90	-0.66	-0.61	-0.49	-1.37	-0.94
A_84_P22154	AT3G29680	AT3G29680	-0.81	-0.59	-1.45	-1.35	-1.87	-1.45	-2.22	-1.92
A_84_P759203	AT3G30730	AT3G30730	-0.59	-0.58	-1.41	-1.29	-1.74	-1.21	-2.16	-1.69
A_84_P286710	AT3G45160	AT3G45160	3.41	3.61	2.20	2.25	2.73	2.85	1.85	2.08
A_84_P526212	AT3G46880	AT3G46880	-1.32	-0.67	-2.62	-2.03	-3.11	-2.68	-3.85	-3.70
A_84_P24099	AT3G47570	AT3G47570	1.91	1.88	0.29	0.15	1.07	0.96	0.26	0.35
A_84_P274520	AT3G47980	AT3G47980	0.51	0.56	-0.55	-0.49	-0.51	-0.39	-1.10	-1.12
A_84_P22211	AT3G48720	AT3G48720	2.61	2.77	0.54	0.37	1.69	1.71	0.23	-0.52
A_84_P24103	AT3G48740	AT3G48740	2.96	1.93	0.59	-0.12	2.13	1.12	-0.53	-1.40
A_84_P813422	AT3G48740	AT3G48740	0.75	-0.15	-1.22	-2.08	0.03	-1.01	-2.45	-2.86
A_84_P593798	AT3G61660	AT3G61660	-1.94	-2.12	-4.01	-3.79	-2.75	-3.18	-4.31	-3.35
A_84_P11867	AT3G62040	AT3G62040	-1.08	-0.40	-3.49	-3.90	-2.18	-2.24	-3.88	-4.19
A_84_P23218	AT3G62930	AT3G62930	1.26	1.45	-0.57	-0.76	-1.05	-0.73	-2.10	-1.80
A_84_P16607	AT4G01450	AT4G01450	1.92	2.16	1.14	1.18	0.30	0.72	-0.93	0.04
A_84_P831869	AT4G02810	AT4G02810	-1.13	-1.70	-2.39	-4.75	-1.72	-2.45	-3.87	-4.18
A_84_P16626	AT4G08770	AT4G08770	1.49	0.71	-2.95	-2.88	-0.19	-1.36	-4.14	-4.85
A_84_P178304	AT4G15690	AT4G15690	0.70	0.13	-1.50	-3.05	-0.49	-0.71	-2.74	-2.70
A_84_P11926	AT4G19380	AT4G19380	3.56	3.53	2.08	2.49	1.72	1.72	1.59	1.92
A_84_P13818	AT4G19820	AT4G19820	0.36	0.12	-2.97	-3.44	-1.65	-1.54	-4.10	-4.81
A_84_P17609	AT4G21870	AT4G21870	5.77	5.36	1.85	1.85	3.09	2.58	1.14	1.60
A_84_P538776	AT4G24015	AT4G24015	0.28	0.61	-0.72	-0.61	-0.67	-0.57	-1.00	-0.72
A_84_P18579	AT4G27300	AT4G27300	2.38	2.44	0.42	0.32	1.74	0.90	-0.51	-1.08
A_84_P151248	AT4G27570	AT4G27570	-1.76	-1.25	-3.30	-4.38	-2.69	-2.17	-3.82	-4.11
A_84_P824302	AT4G29310	AT4G29310	4.25	4.14	2.54	2.48	2.88	2.97	1.99	1.86
A_84_P243195	AT4G29310	AT4G29310	3.66	3.51	1.96	1.82	2.27	2.28	1.34	1.25
A_84_P119612	AT4G31620	AT4G31620	-1.76	-1.20	-2.65	-2.08	-3.35	-2.60	-4.32	-2.58
A_84_P577849	AT4G33770	AT4G33770	2.44	2.54	1.02	0.63	1.71	1.66	0.79	0.51
A_84_P830890	AT4G33770	AT4G33770	2.07	2.16	0.47	0.18	1.42	1.26	0.40	0.17
A_84_P23341	AT4G34790	AT4G34790	1.72	1.71	0.64	0.39	-0.06	0.19	0.16	0.67
A_84_P11048	AT4G34810	AT4G34810	3.21	2.82	0.15	-0.16	-0.01	-0.45	-0.56	-0.30
A_84_P850593	AT4G36040	AT4G36040	5.24	3.99	3.88	3.32	3.84	2.56	3.36	3.04
A_84_P11054	AT4G36040	AT4G36040	4.89	3.59	3.39	2.82	3.42	2.14	2.88	2.61
A_84_P192434	AT4G36850	AT4G36850	5.89	4.15	2.02	0.68	5.10	3.54	1.60	0.20
A_84_P765839	AT4G38825	AT4G38825	1.87	2.05	-2.33	-2.05	1.22	1.35	0.25	0.55
A_84_P16735	AT4G39070	AT4G39070	1.73	2.01	0.72	0.91	0.26	0.24	-0.59	-0.53
A_84_P572544	AT5G03310	AT5G03310	-2.14	-2.45	-3.08	-3.52	-3.33	-3.38	-3.56	-3.29
A_84_P831588	AT5G03360	AT5G03360	-0.06	-0.12	-1.37	-1.73	-0.93	-1.11	-1.04	-1.11

A_84_P732491	AT5G03995	AT5G03995	1.28	1.38	-0.16	-0.28	-2.07	-1.51	-3.17	-2.30
A_84_P237873	AT5G04310	AT5G04310	-0.16	-0.53	-1.80	-1.80	-0.79	-1.48	-2.06	-2.48
A_84_P22447	AT5G05420	AT5G05420	-0.67	-1.31	-2.42	-2.44	-1.58	-2.04	-2.02	-2.54
A_84_P12991	AT5G05790	AT5G05790	0.54	0.39	-0.41	-0.39	-0.74	-0.81	-1.11	-1.56
A_84_P12054	AT5G07580	AT5G07580	4.51	4.57	1.09	1.54	2.34	2.85	0.15	0.65
A_84_P821249	AT5G07580	AT5G07580	2.59	2.69	-0.71	-0.24	0.44	0.97	-1.72	-1.03
A_84_P23593	AT5G08350	AT5G08350	1.74	1.73	-0.79	-0.90	0.32	0.34	-1.21	-1.13
A_84_P311633	AT5G11070	AT5G11070	5.62	5.31	4.49	4.47	3.89	4.34	3.76	4.39
A_84_P597535	AT5G14090	AT5G14090	1.35	1.54	0.00	-0.08	-0.60	-0.31	-1.19	-0.85
A_84_P522030	AT5G14110	AT5G14110	-2.30	-1.93	-3.61	-3.22	-3.28	-3.08	-3.79	-3.73
A_84_P21530	AT5G14120	AT5G14120	5.13	4.73	1.78	1.75	3.56	3.02	1.34	1.35
A_84_P605545	AT5G14330	AT5G14330	0.59	1.61	-0.14	0.66	-0.46	0.82	-0.27	0.59
A_84_P23423	AT5G14360	AT5G14360	-0.16	-0.46	-1.27	-1.44	-1.50	-1.38	-1.11	-1.31
A_84_P309733	AT5G16350	AT5G16350	0.05	0.06	-0.77	-0.95	-0.86	-0.91	-1.09	-1.30
A_84_P11140	AT5G18840	AT5G18840	-0.18	0.50	-1.43	-0.29	-2.70	-2.13	-2.96	-2.28
A_84_P131376	AT5G19190	AT5G19190	3.66	3.01	2.45	2.18	1.56	1.33	1.53	1.62
A_84_P541541	AT5G20640	AT5G20640	-2.70	-1.89	-3.32	-3.29	-3.88	-3.87	-4.09	-3.91
A_84_P795530	AT5G22390	AT5G22390	1.07	0.77	0.31	-0.19	-0.21	-0.56	-0.94	-0.94
A_84_P221276	AT5G22390	AT5G22390	-1.63	-1.84	-2.06	-2.28	-3.42	-3.04	-3.14	-3.14
A_84_P597565	AT5G22870	AT5G22870	-2.01	-2.13	-2.79	-2.87	-3.10	-2.91	-3.37	-3.11
A_84_P10198	AT5G22920	AT5G22920	5.73	5.81	3.89	4.10	3.78	3.90	3.45	3.65
A_84_P814618	AT5G22920	AT5G22920	3.37	3.43	1.69	1.83	1.48	1.59	1.19	1.37
A_84_P10200	AT5G23360	AT5G23360	-1.23	-0.90	-2.31	-2.54	-2.52	-2.18	-3.12	-2.48
A_84_P11146	AT5G23370	AT5G23370	-2.89	-2.79	-4.35	-3.46	-3.94	-3.74	-3.62	-4.21
A_84_P226859	AT5G24570	AT5G24570	3.44	3.04	2.59	2.04	2.66	2.01	2.17	1.79
A_84_P795781	AT5G24570	AK220744	-0.85	-1.13	-1.73	-2.16	-1.23	-1.75	-1.67	-2.04
A_84_P18859	AT5G28630	AT5G28630	1.09	0.95	-0.06	-0.20	0.22	-0.48	-0.22	-0.67
A_84_P502546	AT5G35525	AT5G35525	-1.36	-1.40	-3.91	-3.37	-2.57	-2.32	-3.57	-3.05
A_84_P13065	AT5G39080	AT5G39080	3.81	3.35	2.49	2.25	2.36	2.22	1.74	1.36
A_84_P20636	AT5G40450	AT5G40450	3.06	2.62	0.13	-0.39	2.27	1.85	-0.14	-0.54
A_84_P769282	AT5G40450	AT5G40451	2.03	1.47	-1.06	-1.70	1.22	0.63	-1.16	-1.74
A_84_P56650	AT5G41080	AT5G41080	2.50	2.79	1.76	1.76	0.82	0.60	0.89	1.15
A_84_P790164	AT5G41080	AT5G41080	2.03	2.39	1.41	1.43	0.36	0.33	0.54	0.75
A_84_P594929	AT5G43120	AT5G43120	-2.39	-2.14	-3.46	-2.89	-3.60	-3.57	-4.99	-3.45
A_84_P854040	AT5G44020	AT5G44020	6.45	6.58	5.92	5.98	5.57	5.74	5.81	5.80
A_84_P13082	AT5G44020	AT5G44020	5.14	5.26	4.15	4.28	3.74	3.98	3.44	3.75
A_84_P854498	AT5G44020	AT5G44020	4.59	4.63	3.52	3.70	3.15	3.39	2.86	3.21
A_84_P548712	AT5G49800	AT5G49800	-1.44	-1.17	-2.33	-2.08	-2.07	-1.78	-2.94	-2.57
A_84_P558452	AT5G51850	AT5G51850	-0.93	-0.86	-2.28	-2.04	-2.24	-1.74	-2.28	-2.65
A_84_P5121243	AT5G53230	AT5G53230	1.08	0.26	-0.18	-1.92	0.46	-0.54	-0.21	-2.64
A_84_P159285	AT5G53240	AT5G53240	-1.38	-2.28	-2.83	-4.54	-2.32	-3.51	-2.87	-4.34
A_84_P15958	AT5G54940	AT5G54940	7.00	6.65	6.28	6.05	6.33	5.85	6.06	5.79
A_84_P785967	AT5G54940	AT5G54940	5.65	5.35	5.01	4.79	4.99	4.46	4.75	4.47
A_84_P809682	AT5G54940	AT5G54940	1.21	0.92	0.58	0.40	0.58	-0.01	0.46	0.12
A_84_P11239	AT5G56840	AT5G56840	2.46	2.36	0.08	0.08	1.56	0.79	-1.37	-1.65
A_84_P19829	AT5G58390	AT5G58390	3.73	3.75	1.93	1.89	3.14	3.05	2.30	2.11
A_84_P830800	AT5G58390	AT5G58390	1.45	1.43	-0.39	-0.46	0.90	0.74	-0.07	-0.30
A_84_P12190	AT5G58660	AT5G58660	0.68	0.48	-0.71	-0.93	-0.69	-0.43	-0.10	-0.30
A_84_P834496	AT5G58660	AT5G58660	-1.13	-1.42	-2.59	-2.77	-2.37	-2.41	-2.59	-2.17
A_84_P21648	AT5G58900	AT5G58900	4.00	4.11	1.75	1.93	2.85	3.29	1.44	1.47
A_84_P856315	AT5G60400	AT5G60400	3.76	3.99	2.47	3.08	2.72	3.22	2.31	2.94
A_84_P825265	AT5G60400	AT5G60400	3.78	3.92	2.38	3.01	2.67	3.13	2.26	2.89
A_84_P162513	AT5G60400	AT5G60400	2.15	2.21	0.59	1.21	0.97	1.31	0.34	1.02
A_84_P21661	AT5G61590	AT5G61590	5.82	5.44	0.94	1.68	3.36	3.42	-0.68	0.43
A_84_P17881	AT5G62360	AT5G62360	2.91	3.32	1.33	1.58	0.07	1.80	0.24	1.41
A_84_P847119	AT5G62360	AT5G62360	2.75	3.07	1.17	1.35	-0.17	1.59	0.27	1.29
A_84_P816405	AT5G62360	AT5G62360	0.29	0.66	-1.09	-0.93	-2.69	-0.92	-2.26	-1.01
A_84_P802615	AT5G62360	AT5G62360	-1.32	-0.70	-3.02	-2.48	-4.21	-2.36	-5.00	-2.91
A_84_P297784	AT5G64850	AT5G64850	4.55	4.07	2.64	2.13	3.39	2.48	1.90	1.60
A_84_P243525	AT5G65040	AT5G65040	2.18	2.03	-0.93	-1.19	-0.49	-0.76	-3.37	-3.11
A_84_P571830	AT5G65925	AT5G65925	1.00	1.31	0.15	0.57	0.32	0.58	-0.28	0.39
A_84_P20700	ATL63	AT5G58580	2.81	2.53	1.68	1.46	2.05	1.94	1.13	0.98
A_84_P851410	BE039144		5.97	5.65	4.81	4.95	4.69	3.97	4.02	4.34
A_84_P16748	BES1	AT1G19350	6.71	6.76	6.39	6.35	6.35	6.49	6.52	6.56
A_84_P815715	BES1	AT1G19350	3.68	3.78	3.65	3.62	3.38	3.60	3.67	3.71
A_84_P815721	BES1	AT1G19350	1.28	1.53	0.58	0.59	-0.06	0.26	-0.10	0.18
A_84_P20337	BGAL2	AT3G52840	3.30	3.53	2.44	2.29	2.02	2.35	1.23	1.59
A_84_P12184	BGAL4	AT5G56870	-0.15	-0.66	-0.86	-1.88	-2.03	-1.52	-0.56	-1.32
A_84_P851517	BGAL4	AT5G56870	-2.61	-2.93	-2.83	-3.27	-3.16	-3.71	-2.71	-3.25
A_84_P14235	BGLU46	AT1G61820	-0.81	-0.44	-1.66	-1.81	-2.19	-2.04	-2.43	-2.27
A_84_P17757	BMY3	AT5G18670	5.83	5.41	4.28	4.09	5.21	4.81	4.32	4.17
A_84_P853428	BP803304		-2.05	-1.38	-3.05	-2.69	-3.54	-3.35	-3.88	-3.20
A_84_P228049	BT1	AT5G63160	4.14	2.44	2.38	1.32	2.54	1.57	1.16	0.75
A_84_P21458	BT5	AT4G37610	4.87	4.75	3.60	3.27	3.02	2.75	2.92	2.82

A_84_P16885	BXL1	AT5G49360	3.98	4.28	2.62	2.61	2.08	2.43	1.29	2.25
A_84_P16520	BXL2	AT1G02640	4.63	4.59	2.78	2.60	3.34	3.67	2.95	3.37
A_84_P17164	bZIP	AT1G68880	-1.72	-2.31	-3.36	-3.41	-2.66	-4.17	-4.21	-4.24
A_84_P15068	bZIP3	AT5G15830	-0.49	-0.48	-4.45	-4.48	-2.01	-2.13	-4.39	-4.38
A_84_P17923	BZO2H3	AT5G28770	4.02	3.63	2.12	2.23	2.66	2.49	1.73	2.26
A_84_P822924	BZO2H3	AT5G28770	2.07	1.60	0.14	0.17	0.74	0.56	-0.29	0.29
A_84_P11955	CGA1	AT4G26150	1.44	1.74	-0.26	0.19	0.06	-0.04	-1.90	-1.14
A_84_P22189	CHR38	AT3G42670	1.30	1.85	0.30	0.71	0.45	1.19	-0.14	0.60
A_84_P503591	CIL	AT4G25990	4.12	4.16	1.96	2.31	2.91	2.81	1.28	1.78
A_84_P611792	CKX7	AT5G21482	0.72	0.75	-0.38	-0.05	-0.86	-0.97	-1.04	-1.29
A_84_P265780	CO	AT5G15840	1.56	1.19	-1.05	-1.22	0.02	-0.47	-1.92	-1.41
A_84_P784276	CO	AT5G15840	0.61	0.26	-2.02	-1.88	-0.92	-1.30	-2.45	-1.92
A_84_P584673	CPL3	AT2G33540	1.75	1.73	1.90	2.22	1.71	2.08	1.88	2.20
A_84_P764077	CPL3	AT4G01060	3.02	2.75	1.18	1.99	0.80	1.58	0.38	0.86
A_84_P787734	CPL3	AT4G01060	1.98	1.75	0.21	0.99	-0.18	0.45	-0.54	-0.01
A_84_P532498	CPL3	AT4G01060	2.04	1.82	0.26	1.05	0.00	0.57	-0.59	-0.01
A_84_P14954	CPuORF32	AT1G73600	5.22	5.71	4.18	4.90	2.76	3.68	2.60	3.30
A_84_P10999	CRF2	AT4G23750	4.17	4.27	3.26	3.48	2.83	3.12	3.01	3.49
A_84_P793897	CRF2	AT4G23750	-3.10	-2.41	-3.30	-3.40	-2.84	-3.27	-2.67	-3.13
A_84_P16898	CRF3	AT5G53290	0.92	1.24	-0.34	-0.33	0.26	0.31	-0.75	-0.37
A_84_P15691	CRK30	AT4G11460	2.15	2.14	-1.79	-1.56	-0.91	-0.29	-3.23	-2.63
A_84_P575245	CSLA11	AT5G16190	1.23	1.40	-1.60	-0.25	0.56	-0.30	-2.00	-1.90
A_84_P856597	CSLA11	AT5G16190	1.38	1.51	-1.34	-0.17	-0.24	0.69	-1.81	-1.46
A_84_P16015	CXE17	AT5G16080	0.18	-0.46	-1.08	-1.20	-1.29	-1.68	-1.31	-1.49
A_84_P13313	CYP71B7	AT1G13110	3.18	3.37	1.79	1.96	1.48	1.34	0.87	0.83
A_84_P19565	CYP91A2	AT4G37430	2.45	2.47	1.55	1.73	1.67	1.90	1.15	1.36
A_84_P857969	CYP91A2	AT4G37430	-0.62	-0.51	-1.35	-1.12	-1.22	-1.11	-2.02	-1.78
A_84_P14131	DIN10	AT5G20250	7.25	6.93	3.31	3.20	5.40	4.79	2.44	2.83
A_84_P811412	DIN10	AT5G20250	0.49	0.35	-3.45	-3.28	-1.26	-1.80	-4.13	-3.83
A_84_P856007	DIN10	AT5G20250	-0.43	-0.38	-3.57	-3.46	-1.72	-2.30	-4.56	-4.17
A_84_P811421	DIN10	AT5G20250	-1.45	-1.42	-3.20	-3.74	-2.87	-3.19	-4.12	-3.66
A_84_P853466	DIN10	AT5G20250	-1.00	-1.18	-4.40	-4.21	-2.27	-2.94	-4.64	-4.68
A_84_P795830	DQ108818		-0.49	-1.28	-2.59	-3.92	-1.13	-2.40	-2.47	-3.05
A_84_P797706	EG463912		-2.49	-1.66	-3.20	-2.76	-4.47	-4.19	-4.74	-4.85
A_84_P789279	EG527418		-1.72	-1.56	-2.39	-2.80	-2.85	-2.64	-4.61	-4.05
A_84_P22153	ERD5	AT3G30775	5.50	5.59	1.81	1.69	3.70	2.76	0.76	0.12
A_84_P803506	ERD5	AT3G30775	2.90	3.12	-0.62	-0.65	1.17	0.25	-1.82	-2.34
A_84_P812380	ERD5	AT3G30775	2.30	2.46	-1.26	-1.30	0.63	-0.43	-2.38	-2.89
A_84_P803062	ERD5	AT3G30775	1.99	2.10	-1.51	-1.56	0.28	-0.80	-2.34	-3.22
A_84_P20950	GER1	AT1G72610	7.84	8.07	5.81	6.16	6.92	7.03	5.02	4.90
A_84_P802561	GER1	AT1G72610	6.63	6.93	4.61	4.99	5.75	5.88	3.88	3.75
A_84_P802215	GER1	AT1G72610	-2.98	-2.74	-3.46	-3.20	-2.66	-2.63	-3.99	-3.68
A_84_P19142	GH3.9	AT2G47750	-0.83	-0.95	-1.27	-1.75	-1.77	-1.45	-1.52	-2.16
A_84_P826972	GH3.9	AT2G47750	-3.14	-3.23	-4.07	-4.87	-4.24	-3.99	-4.38	-4.35
A_84_P105796	GRF6	AT2G06200	-0.11	0.19	-3.48	-4.24	-1.29	-0.93	-4.61	-3.90
A_84_P89629	HB22	AT4G24660	2.82	3.06	0.97	1.37	1.82	2.06	0.34	1.01
A_84_P552504	HEXO3	AT1G65590	2.12	2.12	1.14	1.12	1.39	1.55	0.92	0.92
A_84_P822127	HEXO3	AT1G65590	1.85	1.94	0.84	0.81	1.12	1.34	0.66	0.77
A_84_P23171	HSFA7A	AT3G51910	1.42	1.28	0.27	0.11	-0.76	-0.48	-0.41	-0.37
A_84_P16816	IMS2	AT5G23020	-0.79	-1.17	-2.12	-2.59	-2.33	-2.42	-3.80	-3.45
A_84_P186864	IQD12	AT5G03960	-1.49	-1.47	-3.52	-3.63	-3.89	-3.61	-4.36	-4.98
A_84_P581848	JAG	AT1G68480	-1.80	-1.11	-3.55	-2.88	-3.02	-2.62	-4.50	-4.56
A_84_P14981	KAT1	AT5G46240	0.53	0.74	-1.90	-1.83	-0.58	-0.16	-2.66	-2.74
A_84_P18075	KNAT2	AT1G70510	-0.27	0.17	-0.91	-0.77	-1.47	-0.98	-2.79	-1.70
A_84_P22432	LAC8	AT5G01040	-0.94	-0.59	-2.34	-1.49	-3.29	-2.75	-3.66	-2.17
A_84_P277340	LBD25	AT3G27650	-1.69	-1.81	-3.09	-4.16	-3.84	-4.16	-4.37	-4.33
A_84_P57010	LBD39	AT4G37540	5.05	5.04	3.38	3.77	4.09	4.04	2.98	3.23
A_84_P592320	LSH5	AT5G58500	-1.90	-1.50	-2.67	-2.52	-2.95	-3.05	-3.46	-3.31
A_84_P829388	MAPKKK13	AT1G07150	0.46	0.23	-0.73	-1.40	-0.45	-0.87	-0.92	-1.40
A_84_P14478	MES3	AT2G23610	3.36	3.32	1.86	2.04	2.30	2.47	1.33	1.72
A_84_P852772	MES3	AT2G23610	1.30	1.22	-0.22	0.09	0.29	0.48	-0.62	-0.24
A_84_P20466	MIOX4	AT4G26260	-1.36	-0.06	-1.95	-2.47	-3.65	-2.72	-2.96	-2.88
A_84_P784727	MLO10	AT5G65970	1.44	1.42	-0.95	-1.35	0.48	0.32	-1.20	-1.46
A_84_P17895	MLO10	AT5G65970	0.31	0.46	-1.79	-2.46	-0.62	-0.90	-2.58	-2.31
A_84_P15083	MSL9	AT5G19520	-1.57	-0.53	-2.34	-1.29	-2.55	-1.84	-2.21	-1.99
A_84_P11893	MSRB6	AT4G04840	2.71	2.58	1.04	0.87	1.07	1.51	0.70	-0.10
A_84_P15939	MYB111	AT5G49330	0.99	1.53	-0.08	0.63	-0.37	0.34	-0.65	-0.05
A_84_P22464	MYB92	AT5G10280	-2.03	-1.59	-3.13	-3.15	-2.96	-2.68	-4.74	-4.02
A_84_P21604	NAC096	AT5G46590	-1.35	-0.38	-2.85	-2.02	-3.26	-2.39	-3.90	-3.14
A_84_P18118	NAS4	AT1G56430	2.57	3.18	1.82	2.22	1.36	1.55	0.89	1.36
A_84_P15219	NLA	AT1G02860	2.44	2.46	0.57	0.59	1.77	1.63	0.64	0.66
A_84_P839037	NP041809		0.58	0.58	-0.58	-0.84	-0.41	-0.23	-1.39	-1.22
A_84_P11556	NP454661		-1.03	-1.39	-1.75	-2.38	-1.74	-2.24	-2.80	-3.06
A_84_P12427	NRT1:2	AT1G69850	1.47	1.11	-0.27	-0.44	0.83	0.47	-0.54	-0.54

A_84_P821543	NRT1:2	AT1G69850	-0.87	-1.51	-2.67	-2.78	-1.59	-1.97	-2.57	-2.92
A_84_P20458	NRT1.5	AT1G32450	1.58	1.99	0.59	0.61	-1.22	-1.44	-1.95	-1.47
A_84_P587896	OFFP12	AT1G05420	0.72	1.30	-1.97	-1.19	-0.14	0.42	-2.02	-1.39
A_84_P22129	OLEO4	AT3G27660	-0.47	-0.52	-2.61	-3.66	-1.10	-2.90	-3.16	-3.95
A_84_P18581	OPT6	AT4G27730	3.53	3.61	1.10	1.50	1.80	2.21	1.11	1.53
A_84_P11903	ORA59	AT1G06160	1.88	1.80	-0.55	-0.84	0.54	-0.32	-1.02	-1.73
A_84_P10370	PAL3	AT5G04230	1.25	1.15	-1.97	-2.06	0.03	0.32	-2.45	-1.97
A_84_P143149	PDLP3	AT2G33330	4.60	4.76	3.58	3.79	3.61	3.97	2.74	2.93
A_84_P18992	PDR7	AT1G15210	0.21	0.93	-1.64	-0.67	-1.04	-0.57	-2.63	-1.53
A_84_P13563	PK1B	AT2G28930	3.99	4.08	2.43	2.71	3.09	3.44	2.26	2.52
A_84_P789454	PK1B	AT2G28930	3.25	3.35	1.78	1.98	2.40	2.72	1.49	1.83
A_84_P851110	PK1B	AT2G28930	2.90	3.01	1.44	1.69	2.06	2.32	1.02	1.44
A_84_P827013	PKS1	AT2G02950	2.86	2.53	0.59	0.22	1.75	1.65	-0.03	-0.27
A_84_P241573	PKS1	AT2G02950	1.94	1.62	-0.30	-0.57	0.81	0.78	-1.19	-1.22
A_84_P20232	PME3	AT3G14310	8.16	8.16	6.53	7.05	6.66	7.12	6.08	6.46
A_84_P838448	PME3	AT3G14310	4.17	4.18	2.68	3.07	2.78	3.04	2.04	2.37
A_84_P763649	PMEP CRA	AT4G03926	3.85	3.56	2.23	1.95	3.07	2.96	1.86	1.69
A_84_P750624	PMEP CRA	AT1G11580	3.67	3.42	1.91	1.79	2.89	2.73	1.67	1.42
A_84_P846184	PMEP CRA	AT1G11580	1.16	0.87	-0.45	-0.58	0.36	0.32	-0.72	-1.03
A_84_P287770	PP2-A13	AT3G61060	2.28	2.25	1.12	1.35	0.76	0.61	0.60	0.67
A_84_P789365	PP2-A13	AT3G61060	1.67	1.66	0.37	0.73	0.11	0.04	-0.05	0.12
A_84_P11805	PS3	AT3G47420	2.31	2.09	1.60	1.22	1.40	0.96	1.30	1.31
A_84_P142599	PSY1	AT5G58650	2.61	2.29	1.43	0.84	1.88	1.39	1.40	0.92
A_84_P837948	PUP2	AT2G33750	-0.75	-0.60	-2.79	-2.30	-1.41	-1.22	-2.63	-2.62
A_84_P550871	PUP2	AT2G33750	-0.80	-0.65	-3.53	-3.00	-1.43	-1.28	-4.09	-3.77
A_84_P197484	PYL4	AT2G38310	4.80	4.71	1.83	2.27	2.93	2.54	1.01	1.59
A_84_P146199	PYL5	AT5G05440	4.10	3.25	1.86	1.04	2.82	1.98	2.32	2.25
A_84_P817892	PYL5	AT5G05440	2.56	1.81	0.30	-0.47	1.31	0.58	0.84	0.88
A_84_P852047	PYL5	AT5G05440	0.69	-0.06	-1.46	-2.37	-0.61	-1.19	-0.81	-0.93
A_84_P174151	PYL6	AT2G40330	3.38	3.16	-1.17	-1.37	1.62	-0.41	-2.00	-2.54
A_84_P120122	PYR1	AT4G17870	6.09	5.88	3.23	3.45	4.08	3.79	2.78	3.05
A_84_P13235	RAV1	AT1G13260	2.74	2.62	1.66	1.18	1.79	1.89	1.74	1.98
A_84_P94069	RCAR3	AT5G53160	3.42	3.33	1.93	1.86	2.82	2.64	1.41	1.51
A_84_P757994	RL4	AT2G18328	0.81	0.87	-2.47	-2.77	0.14	0.08	-2.75	-3.22
A_84_P262420	RLP26	AT2G33050	1.32	1.38	-0.06	0.07	0.17	0.52	-0.42	-0.04
A_84_P758675	RLP26	AT2G33051	-3.28	-2.88	-3.87	-3.34	-3.88	-3.50	-3.39	-3.95
A_84_P22100	RLP33	AT3G05660	-1.46	-1.92	-2.51	-2.83	-2.28	-2.55	-3.26	-3.19
A_84_P852482	RLP33	AT3G05660	-2.06	-2.51	-3.40	-3.30	-2.93	-3.12	-2.72	-3.41
A_84_P21088	RR16	AT2G40670	2.92	2.99	2.17	1.51	1.88	2.40	1.33	1.55
A_84_P223736	SD1-13	AT1G11350	2.47	2.37	0.95	0.77	1.84	1.78	0.80	0.66
A_84_P868051	SD1-13	AT1G11350	2.15	2.08	0.68	0.51	1.50	1.33	0.45	0.41
A_84_P22400	SEN1	AT4G35770	2.14	0.81	0.96	-0.07	1.18	-0.15	0.32	0.27
A_84_P857167	SEN1	AT4G35770	0.46	-0.58	-0.60	-1.43	-0.29	-1.61	-1.24	-1.29
A_84_P18467	SIP2	AT3G57520	6.18	6.47	5.34	5.61	5.17	4.88	4.77	4.44
A_84_P806643	SIP2	AT3G57520	4.90	5.17	4.20	4.46	3.94	3.65	3.58	3.30
A_84_P597566	SLAH3	AT5G24030	2.14	1.88	1.03	0.41	1.48	0.77	0.56	-0.31
A_84_P576326	SPCH	AT5G53210	2.27	2.29	-0.21	0.27	1.38	1.69	-0.36	0.21
A_84_P14671	SRS6	AT3G54430	1.46	1.63	-0.22	-0.28	0.46	0.84	-1.03	-0.64
A_84_P811601	TA29106_3702		-0.37	-0.45	-1.46	-1.64	-3.12	-2.92	-2.85	-2.66
A_84_P103886	TBL45	AT2G30010	5.75	5.83	3.74	4.00	4.71	4.76	3.55	3.88
A_84_P807831	TBL45	AT2G30010	4.96	5.04	3.05	3.30	3.91	3.99	2.84	3.19
A_84_P850182	TC384346		1.33	1.39	-0.45	-0.11	-1.62	-0.99	-3.20	-1.94
A_84_P101906	TIFY7	AT1G70700	3.79	3.00	2.31	2.37	2.51	2.07	1.74	1.20
A_84_P20456	TLP1	AT4G24180	-1.35	-1.01	-1.93	-1.77	-2.06	-1.50	-2.40	-2.02
A_84_P838419	TLP1	AT4G24180	-2.49	-2.05	-3.38	-3.16	-3.25	-2.81	-3.99	-3.25
A_84_P275730	TMAC2	AT3G02140	3.23	2.84	0.49	0.65	1.93	1.10	-0.05	0.05
A_84_P18147	TPS1	AT1G78580	3.37	3.53	2.61	2.85	2.49	2.58	1.81	1.92
A_84_P850500	TPS1	AT1G78580	-1.76	-1.56	-2.18	-1.99	-2.31	-2.29	-2.39	-2.52
A_84_P20176	UCC2	AT2G44790	-1.14	-0.22	-1.94	-1.35	-2.35	-2.23	-2.44	-2.23
A_84_P19988	UGE1	AT1G12780	3.11	3.10	2.23	2.22	2.31	2.24	1.72	1.72
A_84_P856422	UGE1	AT1G12780	1.16	1.14	0.36	0.26	0.33	0.41	-0.13	-0.10
A_84_P812267	UGE1	AT1G12780	1.01	1.02	0.22	0.07	0.21	0.35	-0.31	-0.42
A_84_P812266	UGE1	AT1G12780	0.93	0.89	0.18	0.11	0.17	0.21	-0.34	-0.34
A_84_P15827	UGT76C2	AT5G05860	3.79	3.53	2.90	2.21	2.78	2.27	2.13	1.56
A_84_P20025	UGT85A2	AT1G22360	4.13	4.03	2.40	2.92	2.62	2.72	1.58	2.11
A_84_P860389	UGT85A2	AT1G22360	-2.03	-1.80	-3.14	-2.82	-3.12	-3.50	-4.97	-4.03
A_84_P17955	UGT85A3	AT1G22380	-1.03	-0.44	-1.74	-1.17	-1.69	-1.38	-2.33	-2.02
A_84_P13341	UGT85A4	AT1G78270	3.12	2.96	1.89	2.14	1.53	2.05	1.45	1.73
A_84_P803356	UGT85A4	AT1G78270	2.74	2.53	1.46	1.74	1.06	1.63	1.02	1.30
A_84_P753447	UGT85A4	AT1G78265	-0.44	-0.59	-0.06	-0.10	-0.12	-0.15	0.00	0.07
A_84_P831400	USPL1	AT1G49320	3.30	3.51	1.48	1.19	2.32	2.74	0.55	1.25
A_84_P147788	USPL1	AT1G49320	-1.20	-1.06	-2.72	-2.98	-1.84	-1.54	-3.77	-3.39
A_84_P232439	WRKY18	AT4G31800	4.58	4.68	1.94	1.92	2.98	2.72	2.08	1.93
A_84_P785249	WRKY18	AT4G31800	2.17	2.30	-0.46	-0.32	0.56	0.27	-0.29	-0.52

A_84_P23291	WRKY29	AT4G23550	1.98	1.95	0.73	1.02	1.08	1.20	0.28	0.81
A_84_P10326	XYL4	AT5G64570	7.59	7.94	5.78	6.12	6.23	6.92	5.10	5.85
A_84_P768803	XYL4	AT5G64572	-1.46	-1.48	-2.41	-1.89	-1.77	-1.96	-2.20	-2.13
A_84_P235383	YSL1	AT4G24120	2.00	1.47	-1.08	-1.16	1.48	0.69	-1.56	-2.13
A_84_P823427	YSL1	AT4G24120	1.67	1.14	-1.19	-1.54	1.08	0.44	-1.66	-2.33
A_84_P16644	YUC2	AT4G13260	0.44	0.22	-2.02	-1.66	-1.34	-0.83	-3.43	-2.74

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