

Supplemental Data S4. Arabidopsis genes utilize APA (APA_class 3) only under salicylic acid treatment

gene	model	tag	total_tags	order_tag	AP1	AP3	AGM	Inflorescence	ROS	SAP	S04	S52	LES	GSE	CAS	SIS	chr_id	strand	position	APA_class
At1g32120	1	GATCTCCATCCAAGACA	44	32	0	0	0	0	0	0	27	0	0	0	0	0	1	w	11554899	3
At1g08720	1	GATCTCGTTATTGCGGA	25	8	0	0	0	0	0	0	25	0	0	0	0	0	1	w	2776774	3
At5g38020	1	GATCTCCCAGACAACGA	8	5	0	0	0	0	0	0	25	0	0	0	0	0	5	c	15184344	3
At3g28430	1	GATCCTGTCACTTCTCT	29	16	0	0	0	0	0	0	21	0	0	0	0	0	3	c	10661553	3
At3g62830	1	GATCTTCAACACCTATG	19	7	0	0	0	0	0	0	18	0	0	0	0	0	3	w	23245064	3
At1g43580	1	GATCTCATGTATAGTGG	8	4	0	0	0	0	0	0	16	0	0	0	0	0	1	c	16413650	3
At3g21160	1	GATCCTGCTGAAGGAAA	21	9	0	0	0	0	0	0	16	0	0	0	0	0	3	c	7415029	3
At4g37920	1	GATCTCTTCTTGAATGA	9	8	0	0	0	0	0	0	16	0	0	0	0	0	4	c	17829862	3
At3g02750	1	GATCAACTGAGGTTGCT	17	10	0	0	0	0	0	0	15	0	0	0	0	0	3	c	595286	3
At3g02750	2	GATCAACTGAGGTTGCT	17	10	0	0	0	0	0	0	15	0	0	0	0	0	3	c	595286	3
At3g02750	3	GATCAACTGAGGTTGCT	14	10	0	0	0	0	0	0	15	0	0	0	0	0	3	c	595286	3
At5g52310	1	GATCCCACCAAAGAAGA	16	11	0	0	0	0	0	0	15	0	0	0	0	0	5	w	21259026	3
At5g66850	1	GATCGCACCTCTTCTGG	19	12	0	0	0	0	0	0	15	0	0	0	0	0	5	c	26715974	3
At4g23600	1	GATCAAGACTTTTGCAA	8	2	0	0	0	0	0	0	14	0	0	0	0	0	4	w	12312531	3
At4g23600	2	GATCAAGACTTTTGCAA	5	2	0	0	0	0	0	0	14	0	0	0	0	0	4	w	12312531	3
At5g61460	1	GATCAAGGATCTTGAAA	22	9	0	0	0	0	0	0	14	0	0	0	0	0	5	c	24734426	3
At2g41220	1	GATCTGATAATCCGGAA	40	23	0	0	0	0	0	0	12	0	0	0	0	0	2	w	17189266	3
At5g55700	1	GATCCGAGAGGTTGGTA	15	13	0	0	0	0	0	0	12	0	0	0	0	0	5	w	22569992	3
At1g48550	1	GATCCACTCAGTGGTGA	13	7	0	0	0	0	0	0	11	0	0	0	0	0	1	c	17955132	3
At5g10180	1	GATCTTGCAACAACAA	19	11	0	0	0	0	0	0	11	0	0	0	0	0	5	w	3194213	3
At1g05030	1	GATCTGAAGTCGAGAAA	14	8	0	0	0	0	0	0	10	0	0	0	0	0	1	c	1439608	3
At1g15570	1	GATCTTCAGTCCAGGAC	19	11	0	0	0	0	0	0	9	0	0	0	0	0	1	w	5363446	3
At2g32430	1	GATCCGAAAACCTCTA	14	11	0	0	0	0	0	0	9	0	0	0	0	0	2	w	13779246	3
At3g63070	1	GATCCGCTCGAACAGAA	22	7	0	0	0	0	0	0	7	0	0	0	0	0	3	w	23318959	3
At5g11610	1	GATCCCACCAAAGCTCA	15	8	0	0	0	0	0	0	7	0	0	0	0	0	5	c	3736690	3
At5g11610	2	GATCCCACCAAAGCTCA	15	8	0	0	0	0	0	0	7	0	0	0	0	0	5	c	3736690	3
At3g46970	1	GATCGTCTCATTCAGCT	22	19	0	0	0	0	0	0	6	0	0	0	0	0	3	c	17316907	3
At2g35450	1	GATCTTCACATTGCAGA	11	6	0	0	0	0	0	0	5	0	0	0	0	0	2	c	14911002	3
At2g17390	1	GATCCAAGTATCAAGGA	17	10	0	0	0	0	0	0	4	0	0	0	0	0	2	w	7563302	3
At3g04500	1	GATCCTACTCTTTCAGA	9	8	0	0	0	0	0	0	3	0	0	0	0	0	3	c	1213122	3
At1g21560	1	GATCATACAGTGTCTTT	9	6	0	0	0	0	0	0	2	0	0	0	0	0	1	w	7555908	3
At2g25620	1	GATCCAGAGAATGGGAG	8	8	0	0	0	0	0	0	2	0	0	0	0	0	2	c	10912032	3
At5g08430	1	GATCAACAATTTAATAA	13	4	0	0	0	0	0	0	2	0	0	0	0	0	5	w	2718983	3
At1g47510	1	GATCTCTTATCCAAAAC	15	8	0	0	0	0	0	0	1	0	0	0	0	0	1	c	17439346	3
At3g57660	1	GATCTTTGGCAGAGTGG	24	22	0	0	0	0	0	0	1	0	0	0	0	0	3	w	21366882	3