Supplementary Table V - Classification of calcium-responsive upregulated genes

Locus	Description	ABRE motif found within promoters - numbers indicate distance in nucleotide upstream of ATG	ABRE-CE motif found within promoters - numbers indicate distance in nucleotides upstream of ATG
1 m 1 G 1 = 500	Transcription factors and DNA/RNA related proteins		
AT4G17500	AtERF-1 - ethylene responsive element binding factor 1	530	
AT3G15210	AtERF4 - ethylene responsive element binding factor 4	234	
AT1G42990	bZIP family transcription factor	317, 158	174
AT1G27730	ZAT10 - salt induced zinc finger protein		
AT1G08930	ERD6 - zinc finger protein ATZF1		
AT2G28840	putative RING zinc finger ankyrin protein		
AT2G40140	CCCH-type zinc finger protein containing ankyrin-repeat		
AT3G55980	PEI1- CCCH zinc finger transcription factor		
AT4G17490	AtERF6 - ethylene-responsive element binding factor 6		186
AT1G18570	myb51 - R2R3 myb like transcription factor		
AT2G30470	transcriptional factor B3 family		
AT3G16720	ATL2 - RING zinc finger protein		
AT3G60220	ATL4 - RING-H2 zinc finger protein	114	
AT4G35480	RING-H2 finger protein RHA3b		
AT3G52800	zinc finger - like protein		
AT3G58510	DEAD BOX PROTEIN 3 (RNA HELICASE-LIKE PROTEIN)		
AT1G66260	export factor RNA binding protein, putative		710
AT3G54770	contains RNA recognition motif		
AT2G45820	remorin - a non-specific DNA binding protein	240	140, 742, 771
	Signaling related proteins		
	calmodulin related		
AT2G41110	ATCAL4		229
AT2G41100	ATCAL5 - TCH3 -CaM related protein	136	
AT5G37770	CALMODULIN-RELATED PROTEIN 2 (TCH2)		124, 143
AT5G42380	calmodulin-related protein	398	
AT3G56800	calmodulin like		335
	Kinases		
AT3G45640	mitogen-activated protein kinase 3		
AT3G59350	protein kinase - Pto kinase interactor 1		933
AT2G01450	ATMPK17 - mitogen-activated protein kinase		
AT1G03740	cdc2-like protein kinase		
AT3G02880	leucine-rich repeat transmembrane protein kinase		
AT1G51805	Similar to light repressible LRR-receptor protein kinase		
AT3G51550	receptor-protein kinase		382, 625
AT2G30740	serine/threonine protein kinase		
AT2G05940	putative protein kinase		
AT3G57530	calcium dependent protein kinase - CDPK		164
	others		
AT1G09070	Similar to Glycine SRC2		900
	cold-regulated protein COR6.6 (KIN2)	128, 146, 219, 440	
AT5G15970	cold-regulated protein COR0.0 (KI112)		
AT5G15970 AT1G20440	COR47 - ERD10		168
		602	168 611
AT1G20440	COR47 - ERD10	602	
AT1G20440 AT2G41430	COR47 - ERD10 ERD15	602	

AT1G76180	dehydrin, putative -ERD14	275, 845	
AT1G47380	protein phosphatase 2C (PPTC)	919	720
AT1G23490	RAB-GTPase - putative ADP-ribosylation factor		
AT5G03530	RAB GTPase		
AT5G62570	calmodulin-binding protein		
AT3G23030	auxin-inducible gene (IAA2)		
AT5G14930	leaf senescence-associated gene SAG101		
AT2G01270	thioredoxin		
AT1G11530	thioredoxin h		
	Transporters		
AT1G59870	ABC transporter family protein	495	359, 375
AT1G59820	similar to Potential phospholipid-transporting ATPase		338, 374, 392, 406
	Photosynthesis		
AT1G55670	putative photosystem I subunit V precursor	493	
AT5G38410	RuBisCO small subunit 3b	218	
	Cytoskeleton and cell structure		
T2G32240	myosin heavy chain -related		147
AT1G21730	kinesin-related protein		
	Defense and PCD related genes		
T2G40000	nematode-resistance protein -related		195
AT5G06320	NDR1/HIN1-like protein 3 (NHL3)		
T5G42500	similar to disease resistance response protein		
T1G33600	LRR protein family		
T1G72910	disease resistance protein (TIR-NBS class), putative		
T1G72930	TIR - TOLL/INTERLEUKIN-1 RECEPTOR-LIKE		
	Proteases		
AT4G02380	ERD1 - SAG21 -clp protease		
AT5G53350	ATP-dependent Clp protease ATP-binding subunit ClpX1	105 151 051	
AT5G59550	COP1-interacting protein CIP8	107, 171, 251	
TDEC 40 (50	Cellular metabolism and others	242	
AT5G42650	AOS - Allene oxide synthase - JA biosynthesis	243	
T1G65310	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like	243	
T1G65310 T1G62660	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase		
AT1G65310 AT1G62660 AT3G55430	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase	243 764	252 328 347 443
AT1G65310 AT1G62660 AT3G55430 AT5G64310	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein		252, 328, 347, 443
AT1G65310 AT1G62660 AT3G55430 AT5G64310 AT1G03220	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP)		252, 328, 347, 443
AT1G65310 AT1G62660 AT3G55430 AT5G64310 AT1G03220 AT1G78850	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family		
AT1G65310 AT1G62660 AT3G55430 AT5G64310 AT1G03220 AT1G78850 AT2G22500	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family		341
AT1G65310 AT1G62660 AT3G55430 AT5G64310 AT1G03220 AT1G78850 AT2G22500 AT4G24570	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family		
T1G65310 T1G62660 T3G55430 T5G64310 T1G03220 T1G78850 T2G22500 T4G24570 T2G30870	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family mitochondrial carrier protein family GST - ERD13		341
T1G65310 T1G62660 T3G55430 T5G64310 T1G03220 T1G78850 T2G22500 T4G24570 T2G30870 T1G27130	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family mitochondrial carrier protein family GST - ERD13 GST12		341
T1G65310 T1G62660 T3G55430 T5G64310 T1G03220 T1G78850 T2G22500 T4G24570 T2G30870 T1G27130 T1G76680	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family mitochondrial carrier protein family GST - ERD13 GST12	764	341
T1G65310 T1G62660 T3G55430 T5G64310 T1G03220 T1G78850 T2G22500 T4G24570 T2G30870 T1G27130 T1G76680 T5G56870	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family mitochondrial carrier protein family GST - ERD13 GST12 12-oxophytodienoate reductase (OPR1) glycosyl hydrolase family 35 (beta-galactosidase)	764 893	341 228
T1G65310 T1G62660 T3G55430 T5G64310 T1G03220 T1G78850 T2G22500 T4G24570 T2G30870 T1G27130 T1G76680 T1G76680 T1G76680	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family mitochondrial carrier protein family GST - ERD13 GST12 12-oxophytodienoate reductase (OPR1) glycosyl hydrolase family 35 (beta-galactosidase) glycosyltransferase family 8	764	341
T1G65310 T1G62660 T3G55430 T5G64310 T1G03220 T1G78850 T2G22500	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family mitochondrial carrier protein family GST - ERD13 GST12 12-oxophytodienoate reductase (OPR1) glycosyl hydrolase family 35 (beta-galactosidase) glycosyltransferase family 8 UDP-glucose glucosyltransferase	764 893	341 228
T1G65310 T1G62660 T3G55430 T5G64310 T1G03220 T1G78850 T2G22500 T4G24570 T2G30870 T1G27130 T1G76680 T1G76680 T1G76680 T1G76090 T1G22360	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family mitochondrial carrier protein family GST - ERD13 GST12 12-oxophytodienoate reductase (OPR1) glycosyl hydrolase family 35 (beta-galactosidase) glycosyltransferase family 8 UDP-glucose glucosyltransferase UDP-glucose dehydrogenase -related	764 893	341 228 15, 201, 219, 935
T1G65310 T1G62660 T3G55430 T5G64310 T1G03220 T1G78850 T2G22500 T4G24570 T2G30870 T1G27130 T1G76680 T5G56870 T1G70090 T1G22360 T3G29360	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family mitochondrial carrier protein family GST - ERD13 GST12 12-oxophytodienoate reductase (OPR1) glycosyl hydrolase family 8 UDP-glucose glucosyltransferase UDP-glucose dehydrogenase -related GDP-mannose pyrophosphorylase	764 893	341 228 15, 201, 219, 935
T1G65310 T1G62660 T3G55430 T5G64310 T1G03220 T1G78850 T2G22500 T4G24570 T2G30870 T1G27130 T1G76680 T5G56870 T1G70090 T1G22360 T3G29360 T2G39770	AOS - Allene oxide synthase - JA biosynthesis xyloglucan endotransglycosylase TCH4-like glycosyl hydrolase family 32, identical to beta-fructosidase glycosyl hydrolase family 17 -beta 1,3 glucanase AGP1 - arabinogalactan-protein extracellular dermal glycoprotein (EDGP) curculin-like (mannose-binding) lectin family mitochondrial carrier protein family mitochondrial carrier protein family GST - ERD13 GST12 12-oxophytodienoate reductase (OPR1) glycosyl hydrolase family 35 (beta-galactosidase) glycosyltransferase family 8 UDP-glucose glucosyltransferase UDP-glucose dehydrogenase -related	764 893	341 228 15, 201, 219, 935

AT2G46330	AGP16 - arabinogalactan-protein (AGP16)		
AT5G13420	transaldolase - like protein		
AT1G55810	uracil phosphoribosyltransferase		
AT4G39940	adenosine-5-phosphosulfate-kinase		
AT1G73540	Similar to diphosphoinositol polyphosphate phosphohydrolase	641	176, 504, 549
AT1G74100	putative flavonol sulfotransferase	457	
AT2G45300	5-enolpyruvylshikimate-3-phosphate (EPSP) synthase	263	950
AT4G33070	pyruvate decarboxylase-1 (Pdc1)		
AT5G11670	NADP dependent malic enzyme - like protein	221	
AT5G25880	malate dehydrogenase		
AT3G17390	MTO3 - s-adenosylmethionine synthetase - Met biosynthesis		498
AT5G05730	ASA1 - anthralinate synthase alpha subunit - Trp biosynthesis		
AT3G44720	prephenate dehydratase family, similar to bacterial PheA		
AT4G30530	component of aniline dioxygenase	155	
AT4G38550	Phospholipase like protein-pEARLI 4		200
AT5G24140	squalene monooxygenase 2	278	
AT1G21110	O-methyltransferase 1, putative		
AT2G05710	cytoplasmic aconitate hydratase	647, 932	
AT5G26030	ferrochelatase-I		
AT4G22710	CYP706A2 - cytochrome p450		295
AT5G57220	CYP81F2 cytochrome p450		
AT4G31500	CYP83B1, CYTOCHROME P450 MONOOXYGENASE		
AT4G39950	cytochrome P450		
AT3G15356	similar to putative lectin		
AT3G16530	putative lectin		
AT1G79900	mitochondrial carrier protein family		
AT3G11820	SYP121 - syntaxin-related protein		
AT5G13650	GTP-binding protein typA (tyrosine phosphorylated protein A)		
AT2G29960	cyclophilin		
AT3G19260	LAG1 HOMOLOG 2, LONGEVITY ASSURANCE GENE1		
AT3G26980	similar to geranylgeranylated proteins ATGP4		554
AT5G03380	similar to farnesylated protein ATFP2	784	
	unknown genes		
AT2G23120	unknown protein		
AT1G19180	unknown protein	284, 342	666, 816
AT1G30360	unknown protein		
AT1G69890	unknown protein		
AT2G41640	unknown protein		
AT4G39670	unknown protein		
AT5G42050	unknown protein		658
AT1G03730	unknown protein		213, 257
AT1G18470	unknown protein		287
AT1G28380	unknown protein		
AT1G31130	unknown protein		
AT1G36310	unknown protein		
AT1G36580	unknown protein	151	
AT1G49480	unknown protein		
AT1G52890	unknown protein		795, 858
AT1G62870	unknown protein		375, 359
AT1G69080	unknown protein		253
AT1G69840	unknown protein		
AT1G72450	unknown protein	904, 941	
AT1G74950	unknown protein	255, 860	
AT2G01670	unknown protein		
AT2G16530	unknown protein		

AT2G20240	unknown protein		
AT2G20810	unknown protein		
AT2G23810	unknown protein	525	387
At2G26190	unknown protein		
AT3G05320	unknown protein		
AT3G07180	unknown protein	171	155
AT3G07780	unknown protein		755
AT3G17650	unknown protein	545	
AT3G29170	unknown protein		311
AT3G56880	unknown protein		204, 234
AT3G62680	proline-rich protein		
AT4G21850	unknown protein		
AT4G24160	unknown protein		178
AT4G25030	unknown protein	676	793
AT4G32020	unknown protein		252
AT4G36500	unknown protein		
AT5G03210	unknown protein	351	175, 539
AT5G09850	unknown protein		
AT5G18400	unknown protein		499
AT5G45350	unknown protein		
AT5G46500	unknown protein		
AT5G56980	unknown protein	232	
AT5G65630	unknown protein	162, 431	