Appendix 2c Genes up-regulated following the freezing treatment. This table contains the full set of genes that were responsive to the freezing treatment (P <0.05) and that were up-regulated (log2 fold change in expression >-1).

	Fold		
	change		
Probe ID	(log2)	AGI Code	TAIR annotation
			similar to unknown protein [Arabidopsis thaliana] (TAIR:AT5G12010.1); similar to
			unnamed protein product [Vitis vinifera] (GB:CAO43835.1); contains domain
	3.5226	At4g29780	PTHR22930 (PTHR22930)
266545_at	3.1542	At2g35290	similar to unnamed protein product [Vitis vinifera] (GB:CAO63442.1)
			TCH4_XTH22TCH4 (TOUCH 4); hydrolase, acting on glycosyl bonds /
247925_at	3.0612	At5g57560	xyloglucan:xyloglucosyl transferase
			similar to unknown protein [Arabidopsis thaliana] (TAIR:AT1G18740.1); similar to
			unnamed protein product [Vitis vinifera] (GB:CAO43630.1); contains InterPro
260227_at	3.0470	At1g74450	domain Protein of unknown function DUF793 (InterPro:IPR008511)
255064_at	2.8735	At4g08950	phosphate-responsive protein, putative (EXO)
			STZ_ZAT10STZ (SALT TOLERANCE ZINC FINGER); nucleic acid binding /
261648_at	2.7731	At1g27730	transcription factor/ zinc ion binding
		At2g05050	
		At2g30020	
000004	0.005	At3g27140	
266834_s_at	2.6695	At4g08260	laimillar to unknown protein [Archidencie theliane] (TAID: AT1C74450 4); similar to
			similar to unknown protein [Arabidopsis thaliana] (TAIR:AT1G74450.1); similar to
004405 64	0.5040	A+1 ~10710	unnamed protein product [Vitis vinifera] (GB:CAO43630.1); contains InterProdomain Protein of unknown function DUF793 (InterPro:IPR008511)
261405_at	2.5616	At1g18740	GATL10GATL10 (Galacturonosyltransferase-like 10); polygalacturonate 4-alpha-
256633_at	2.4828	At3g28340	galacturonosyltransferase/ transferase, transferring hexosyl groups
262307_at	2.3694	At1g71000	DNAJ heat shock N-terminal domain-containing protein
254120_at	2.3304	At4g24570	mitochondrial substrate carrier family protein
257654_at	2.2573	At3g13310	DNAJ heat shock N-terminal domain-containing protein
257034_at	2.1959	At3g46620	zinc finger (C3HC4-type RING finger) family protein
202474_at	2.1303	7110g-10020	CBF4_DREB1DCBF4/DREB1D (C- REPEAT-BINDING FACTOR 4); DNA
248389_at	2.0986	At5g51990	binding / transcription activator/ transcription factor
253859_at	2.0907	At4g27657	similar to unknown protein [Arabidopsis thaliana] (TAIR:AT4G27652.1)
253830_at	2.0727	At4g27652	similar to unknown protein [Arabidopsis thaliana] (TAIR:AT4G27657.1)
		At2g05440	[
265561_s_at	2.0214	At2g05510	glycine-rich protein
252679_at	1.9511	At3g44260	CCR4-NOT transcription complex protein, putative
		J	ATERF11_ERF11_ATERF11/ERF11 (ERF domain protein 11); DNA binding /
261470_at	1.9067	At1g28370	transcription factor/ transcription repressor
253915_at	1.8822	At4g27280	calcium-binding EF hand family protein
246011_at	1.8594	At5g08330	TCP family transcription factor, putative
247478_at	1.8330	At5g62360	invertase/pectin methylesterase inhibitor family protein
247177_at	1.7487	At5g65300	unknown protein
259788_at	1.7231	At1g29670	GDSL-motif lipase/hydrolase family protein
		At1g68050	
		At5g23410	FKF1_ADO3FKF1 (FLAVIN-BINDING KELCH DOMAIN F BOX PROTEIN);
259990_s_at		At5g42730	ubiquitin-protein ligase
264005_at	1.6282	At2g22470	AGP2_AGP2 (ARABINOGALACTAN-PROTEIN 2)
266037_at	1.6116	At2g05940	protein kinase, putative
251640_at	1.5780	At3g57450	similar to unnamed protein product [Vitis vinifera] (GB:CAO40798.1)
			MAPKKK14MAPKKK14 (Mitogen-activated protein kinase kinase kinase 14);
266832_at	1.5544	At2g30040	kinase
266123_at	1.5429	At2g45180	protease inhibitor/seed storage/lipid transfer protein (LTP) family protein
267411_at	1.5050	At2g34930	disease resistance family protein
254667_at	1.4985	At4g18280	glycine-rich cell wall protein-related
248243_at	1.4816	At5g53590	auxin-responsive family protein

252475 s. at 1.3493	253104_at	1.4431	At4g36010	pathogenesis-related thaumatin family protein
Aliga6664   Aliga9570   Aliga9670   Aliga970				
252475 s. at 1.3436	200070_at	1.0070		1412014120 (142141) III (142141)
248966_at   1.3298	252475 s at	1.3436		LUX PCL1 PCL1 (PHYTOCLOCK 1): DNA binding
248146 at   1.3067				
248964_at   1.2731				
288964_at   1.2731				
248964_at         1.2731         At5g45340_3): oxygen binding           255926_at         1.2709         At1g22190         AP2 domain-containing transcription factor, putative           255926_at         1.2709         At1g22190         AP2 domain-containing transcription factor, putative           similar to unknown protein [Arabidopsis thaliana] (TAIR:AT1G64650.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT1G64650.1); similar to unknown protein product [Vitis vinifera] (GB:CA062932.1); contains InterPro domain Protein of unknown function DUF791 (InterPro!PR06309); contains InterPro domain MFS general substrate transport (InterPro!PR06809); contains InterPro domain MFS general substrate transport (InterPro!PR016196)           252377_at         1.2627         At3g47960         proton-dependent oligopeptide transport (POT) family protein           2526937_at         1.2244         At2g17660         nitrate-responsive NOI protein, putative           256093_at         1.2290         At3g56880         VO motif-containing protein           247280_at         1.1584         At3g15210         protein binding / transcription factor / transcription repressor           251861_at         1.1571         At3g54810         BME3_BME33_ME33_F (BLUE MICROPYLAR END3); transcription factor           247280_at         1.1494         At5g64260         phosphate-responsive protein, putative           258903_at         1.1370         At3g29030         At3g29303			i i i ge e e e	
255926_at	248964_at	1.2731	At5g45340	
protein [Arabidopsis thaliana] (TARR.AT3G49310.1); similar to unknown [Populus trichocarpa] (GB:ABK93582.1); similar to unnamed protein product [Vitis vinifera] (GB:CAO48607.1); similar to unnamed protein product [Vitis vinifera] (GB:CAO48607.1); similar to unnamed protein product [Vitis vinifera] (GB:CAO62932.1); contains InterPro domain Protein of unknown function DUF791 (InterPro:IPR008509); contains InterPro domain Protein of unknown function DUF791 (InterPro:IPR008509); contains InterPro domain MFS general substrate transport and protein product [Vitis vinifera] (GB:CAO62932.1); contains InterPro domain Protein of unknown function DUF791 (InterPro:IPR016196)  25377 at 1.2627				similar to unknown protein [Arabidopsis thaliana] (TAIR:AT1G64650.2); similar to
trichocarpa   (GB:CAO48677.1); similar to unnamed protein product [Vitis vinifera] (GB:CAO48677.1); similar to unnamed protein product [Vitis vinifera] (GB:CAO62932.1); contains InterPro domain Protein of unknown function DUF791 (InterPro:IPR008509); contains InterPro domain MFS general substrate transport and the provided protein product [Vitis vinifera] (InterPro:IPR008509); contains InterPro domain MFS general substrate transport and the protein product [Vitis vinifera] (InterPro:IPR008509); contains InterPro domain MFS general substrate transport and protein protein and protein and protein protein protein and protein protein and protein p				unknown protein [Arabidopsis thaliana] (TAIR:AT1G64650.1); similar to unknown
GB:CAO48607.1); similar to unnamed protein product [Vitis vinifera] (GB:CAO62932.1); contains InterPro domain Protein of unknown function DUF791 (InterPro:IPR008509); contains InterPro domain MFS general substrate transporter (InterPro:IPR016196)				protein [Arabidopsis thaliana] (TAIR:AT3G49310.1); similar to unknown [Populus
CB:CAO62932.1); contains InterPro domain Protein of unknown function DUF791 (InterPro:IPR008509); contains InterPro domain MFS general substrate transporter (InterPro:IPR008509); contains InterPro domain MFS general substrate transport (InterPro:IPR016196)				trichocarpa] (GB:ABK93582.1); similar to unnamed protein product [Vitis vinifera]
1.2681				(GB:CAO48607.1); similar to unnamed protein product [Vitis vinifera]
253891_at         1.2681         At4g27720         transporter (InterPro:IPR016196)           252377_at         1.2627         At3g47860         proton-dependent oligopeptide transport (POT) family protein           256093_at         1.2290         At1g20823         zinc finger (C3HC4-type RING finger) family protein           246289_at         1.2229         At3g56880         VQ motif-containing protein           257053_at         1.1584         At3g15210         protein binding / transcription factor/ transcription repressor           257053_at         1.1584         At3g15210         protein binding / transcription factor/ transcription repressor           257053_at         1.1571         At3g54810         BME3_BME3/BME3-ZF (BLUE MICROPYLAR END3); transcription factor           247280_at         1.1494         At5g64260         phosphate-responsive protein, putative           258003_at         1.1370         At3g29303         THALIANA EXPANSIN A5)           258003_at         1.0953         At5g63130         octicosapeptide/Phox/Bem1p (PB1) domain-containing protein           263769_at         1.0991         At2g24550         at4g24550           263799_at         1.0914         At5g22630         prophenate dehydratase           264000_at         1.0914         At5g22630         prophenate dehydratase				(GB:CAO62932.1); contains InterPro domain Protein of unknown function DUF791
252377_at         1.2627         At3g47960         proton-dependent oligopeptide transport (POT) family protein           264617_at         1.2441         At2g17660         nitrate-responsive NOI protein, putative           256093_at         1.2290         At3g56880         VQ motif-containing protein           246289_at         1.2291         At3g56880         VQ motif-containing protein           ATERF4_ATERF4_ERF4_RAP2.5_ATERF-4/ATERF4/ERF4/RAP2.5         (ETHYLENE RESPONSIVE ELEMENT BINDING FACTOR 4); DNA binding / transcription factor/ transcription repressor           257053_at         1.1584         At3g15210         protein binding / transcription factor/ transcription repressor           257861_at         1.1571         At3g54810         BME3_BME3/BME3-ZF (BLUE MICROPYLAR END3); transcription factor           247280_at         1.1494         At5g64260         phosphate-responsive protein, putative           258003_at         1.1370         At3g29030         THALIANA EXPANSIN A5)           263662_at         1.1241         At1g04430         dehydration-responsive protein-related           247393_at         1.0953         At2g63130         octicosapeptide/Phox/Bem1p (PB1) domain-containing protein           263799_at         1.0951         At2g24550         unknown [Populus trichocarpa x Populus deltoides] (GB:ABK96563.1)           249910_at         1.0954				(InterPro:IPR008509); contains InterPro domain MFS general substrate
252377_at   1.2627	253891_at	1.2681	At4g27720	transporter (InterPro:IPR016196)
264617_at         1.2441         At2g17660         nitrate-responsive NOI protein, putative           256093_at         1.2299         At1g20823         zinc finger (C3HC4-type RING finger) family protein           246289_at         1.2299         At3g56880         VQ motif-containing protein           ATERF4_ATERF4_ERF4_RAP2.5         ATERF4/ATERF4/ERF4/RAP2.5           (ETHYLENE RESPONSIVE ELEMENT BINDING FACTOR 4); DNA binding / protein binding / transcription factor/ transcription repressor           257053_at         1.1584         At3g54810         BME3_BME3/BME3-2F (BLUE MICROPYLAR END3); transcription factor           247280_at         1.1494         At5g64260         phosphate-responsive protein, putative           ATEXPA5_ATEXP5_ATHEXP ALPHA 1.4ATEXPA5 (ARABIDOPSIS         ATEXPA5_ATEXP5_ATHEXP ALPHA 1.4ATEXPA5 (ARABIDOPSIS           258003_at         1.1370         At3g29030         THALIANA EXPANSIN A5)           263662_at         1.1241         At1g04430         dehydration-responsive protein-related           247393_at         1.0953         At5g63130         octicosapeptide/Phox/Bem1p (PB1) domain-containing protein           263799_at         1.0951         At2g24550         mitochondrial substrate carrier family protein           249910_at         1.0914         At5g22630         mitochondrial substrate carrier family protein           261318_at			At3g47960	proton-dependent oligopeptide transport (POT) family protein
256093_at   1.229		1.2441	At2g17660	
246289_at         1.2229         At3g56880         VQ motif-containing protein           ATERF4_ATERF4_ERF4_RAP2.5         ATERF4_ATERF4_ERF4_RAP2.5 (ETHYLENE RESPONSIVE ELEMENT BINDING FACTOR 4); DNA binding / protein binding / transcription factor / transcription repressor           257053_at         1.1584         At3g15210         protein binding / transcription factor / transcription repressor           251861_at         1.1571         At3g54810         BME3_BME3/BME3-ZF (BLUE MICROPYLAR END3); transcription factor           247280_at         1.1494         At5g64260         phosphate-responsive protein, putative           ATEXPAS_ATEXPS_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS)         ATEXPAS_ATEXPS_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS)           258003_at         1.1370         At3g29030         THALIANA EXPANSIN A5)           263662_at         1.1241         At1g04430         dehydration-responsive protein-related           247393_at         1.0953         At5g63130         octicosapeptide/Phox/Bem1p (PB1) domain-containing protein           263799_at         1.0914         At5g224550         unknown protein [Arabidopsis thaliana] (TAIR:AT4G31510.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to unknown [Populus trichocarpa] (GB:ABK95652.1)           250098				zinc finger (C3HC4-type RING finger) family protein
ATERF4_ATERF-4_ERF4_RAP2.5_ATERF-4/ATERF4/ERF4/RAP2.5 (ETHYLENE RESPONSIVE ELEMENT BINDING FACTOR 4); DNA binding / protein binding / transcription factor/ transcription repressor 251861_at 1.1571 At3g54810 BME3_BME3/BME3-ZF (BLUE MICROPYLAR END3); transcription factor 47280_at 1.1494 At5g64260 phosphate-responsive protein, putative ATEXPA5_ATEXPA5_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS ATMEXPA5_ATEXPA5_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS ATMEXPA3_ATEXPA5_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS ATMEXPA5_ATEXPA5_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS ATMEXPA5_ATEXPA5_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS ATMEXPA5_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS ATMEXPA5_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS ATMEXPA5_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS ATMEXPA5_ATHEXP ALPHA 1.4_ATEXPA5_ATHEXP ALPHA				
(ETHYLENE RESPONSIVE ELEMENT BINDING FACTOR 4); DNA binding / protein binding / protein binding / transcription factor/ transcription repressor	_		J	
251861_at         1.1571         At3g54810         BME3_BME3_BME3/BME3-ZF (BLUE MICROPYLAR END3); transcription factor           247280_at         1.1494         At5g64260         phosphate-responsive protein, putative           258003_at         1.1370         At3g29030         THALIANA EXPANSIN A5)           263662_at         1.1241         At1g04430         dehydration-responsive protein-related           247393_at         1.0953         At5g63130         octicosapeptide/Phox/Bem1p (PB1) domain-containing protein           263799_at         1.0951         At2g24550         acticosapeptide/Phox/Bem1p (PB1) domain-containing protein           249910_at         1.0914         At2g24550         milothondrial substrate carrier family protein           249910_at         1.0903         At2g22500         mitochondrial substrate carrier family protein           261318_at         1.0876         At1g53035         mitochondrial substrate carrier family protein           250098_at         1.0841         At5g17350         unknown [Populus trichocarpa] (GB:ABK95625.1)           261285_at         1.0669         At1g75800         pathogenesis-related thaumatin family protein           253286_at         1.0617         At3g05910         pectinacetylesterase, putative           253286_at         1.0410         At5g49330         AtMYB111_AMYB111 (myb domai				
251861_at         1.1571         At3g54810         BME3_BME3_BME3/BME3-ZF (BLUE MICROPYLAR END3); transcription factor           247280_at         1.1494         At5g64260         phosphate-responsive protein, putative           258003_at         1.1370         At3g29030         THALIANA EXPANSIN A5)           263662_at         1.1241         At1g04430         dehydration-responsive protein-related           247393_at         1.0953         At5g63130         octicosapeptide/Phox/Bem1p (PB1) domain-containing protein           263799_at         1.0951         At2g24550         acticosapeptide/Phox/Bem1p (PB1) domain-containing protein           249910_at         1.0914         At2g24550         milothondrial substrate carrier family protein           249910_at         1.0903         At2g22500         mitochondrial substrate carrier family protein           261318_at         1.0876         At1g53035         mitochondrial substrate carrier family protein           250098_at         1.0841         At5g17350         unknown [Populus trichocarpa] (GB:ABK95625.1)           261285_at         1.0669         At1g75800         pathogenesis-related thaumatin family protein           253286_at         1.0617         At3g05910         pectinacetylesterase, putative           253286_at         1.0410         At5g49330         AtMYB111_AMYB111 (myb domai	257053 at	1.1584	At3q15210	protein binding / transcription factor/ transcription repressor
247280_at         1.1494         At5g64260         phosphate-responsive protein, putative           258003_at         1.1370         At3g29030         THALIANA EXPANSIN A5)           263662_at         1.1241         At1g04430         dehydration-responsive protein-related           247393_at         1.0953         At5g63130         octicosapeptide/Phox/Bem1p (PB1) domain-containing protein           263799_at         1.0951         At2g24550         unknown protein [Arabidopsis thaliana] (TAIR:AT4G31510.1); similar to unknown [Populus trichocarpa x Populus deltoides] (GB:ABK96563.1)           249910_at         1.0914         At5g22630         prephenate dehydratase           264000_at         1.0903         At2g22500         mitochondrial substrate carrier family protein           36739_at         1.0876         At1g53035         unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to unknown protein [Arabidopsis thaliana]           250098_at         1.0841         At1g75800         At1g75800<		1		
ATEXPA5_ATEXP5_ATHEXP ALPHA 1.4_ATEXPA5 (ARABIDOPSIS 1.1370 At3g29030 THALIANA EXPANSIN A5)  263662_at 1.1241 At1g04430 dehydration-responsive protein-related 247393_at 1.0953 At5g63130 octicosapeptide/Phox/Bem1p (PB1) domain-containing protein similar to unknown protein [Arabidopsis thaliana] (TAIR:AT4G31510.1); similar to unknown [Populus trichocarpa x Populus deltoides] (GB:ABK96563.1) ADT5_ADT5 (AROGENATE DEHYDRATASE 5); arogenate dehydratase/ prephenate dehydratase 264000_at 1.0903 At2g22500 mitochondrial substrate carrier family protein similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to unknown [Populus trichocarpa] (GB:ABK95625.1) at 1.0669 At1g75800 pathogenesis-related thaumatin family protein ANNAT1_ATOXY5_OXY5_ANNAT1 (ANNEXIN ARABIDOPSIS 1); calcium ion binding / calcium-dependent phospholipid binding bectinacetylesterase, putative 253286_at 1.0410 At3g05910 pectinacetylesterase, putative 247655_at 1.0351 At4g34260 RHA3B_RHA3B_RHNG-H2 finger A3B); protein binding / transcription RHL41_ZAT12_RHL41 (RESPONSIVE TO HIGH LIGHT 41); nucleic acid binding / transcription factor/ zinc ion binding / MRH5_SHV3_MRH5/SHV3 (morphogenesis of root hair 5);				
258003_at         1.1370         At3g29030         THALIANA EXPANSIN A5)           263662_at         1.1241         At1g04430         dehydration-responsive protein-related           247393_at         1.0953         At5g63130         octicosapeptide/Phox/Bem1p (PB1) domain-containing protein           263799_at         1.0951         At2g24550         unknown [Populus trichocarpa x Populus deltoides] (GB:ABK96563.1)           ADT5_ADT5 (AROGENATE DEHYDRATASE 5); arogenate dehydratase/         At5g22630         prephenate dehydratase           249910_at         1.0903         At2g22500         mitochondrial substrate carrier family protein           261318_at         1.0876         At1g53035         mitochondrial substrate carrier family protein           250098_at         1.0841         At5g17350         unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3	_		J	
263662_at         1.1241         Attg04430         dehydration-responsive protein-related           247393_at         1.0953         At5g63130         octicosapeptide/Phox/Bem1p (PB1) domain-containing protein           263799_at         1.0951         At2g24550         unknown protein [Arabidopsis thaliana] (TAIR:AT4G31510.1); similar to unknown [Populus trichocarpa x Populus deltoides] (GB:ABK96563.1)           249910_at         1.0914         At5g22630         prephenate dehydratase           264000_at         1.0903         At2g22500         mitochondrial substrate carrier family protein           261318_at         1.0876         At1g53035         unnamed protein product [Vitis vinifera] (GB:CAO69750.1)           250098_at         1.0841         At5g17350         unknown [Populus trichocarpa] (GB:ABK95625.1)           262727_at         1.0669         At1g75800         pathogenesis-related thaumatin family protein           261285_at         1.0648         At1g35720         pathogenesis-related thaumatin family protein           253286_at         1.0555         At4g34260         catalytic           248596_at         1.0410         At5g49330         AtWB111_AtMYB111 (myb domain protein 111); DNA binding / transcription           253140_at         1.0384         At4g35480         RHA3B_RHA3B (RING-H2 finger A3B); protein binding / zinc ion binding           247655	258003 at	1.1370	At3g29030	
247393_at 1.0953 At5g63130 octicosapeptide/Phox/Bem1p (PB1) domain-containing protein similar to unknown protein [Arabidopsis thaliana] (TAIR:AT4G31510.1); similar to unknown [Populus trichocarpa x Populus deltoides] (GB:ABK96563.1)  ADT5_ADT5 (AROGENATE DEHYDRATASE 5); arogenate dehydratase/pephenate dehydratase arrier family protein  249910_at 1.0903 At2g22500 mitochondrial substrate carrier family protein  261318_at 1.0876 At1g53035 unnamed protein product [Vitis vinifera] (GB:CAO69750.1)  similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to unknown [Populus trichocarpa] (GB:ABK95625.1)  250098_at 1.0841 At5g17350 pathogenesis-related thaumatin family protein  ANNAT1_ATOXY5_OXY5ANNAT1 (ANNEXIN ARABIDOPSIS 1); calcium ion binding / calcium-dependent phospholipid binding  251286_at 1.0648 At1g35720 binding / calcium-dependent phospholipid binding  253286_at 1.0410 At5g49330 AtMYB111_AtMYB111 (myb domain protein 111); DNA binding / transcription  253140_at 1.0384 At4g35480 RHA3B_RHA3B (RING-H2 finger A3B); protein binding / zinc ion binding  RHL41_ZAT12_RHL41 (RESPONSIVE TO HIGH LIGHT 41); nucleic acid binding / transcription factor/ zinc ion binding  MRH5_SHV3_MRH5/SHV3 (morphogenesis of root hair 5);			At1g04430	dehydration-responsive protein-related
similar to unknown protein [Arabidopsis thaliana] (TAIR:AT4G31510.1); similar to unknown [Populus trichocarpa x Populus deltoides] (GB:ABK96563.1)  At5g22630 prephenate dehydratase  249910_at 1.0914 At5g22630 prephenate dehydratase  264000_at 1.0903 At2g22500 mitochondrial substrate carrier family protein  Similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unnamed protein product [Vitis vinifera] (GB:CAO69750.1)  Similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unnamed protein product [Vitis vinifera] (GB:CAO69750.1)  Similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unknown [Populus trichocarpa] (GB:ABK95625.1)  250098_at 1.0841 At5g17350 pathogenesis-related thaumatin family protein  ANNAT1_AT0XY5_OXY5_ANNAT1 (ANNEXIN ARABIDOPSIS 1); calcium ion binding / calcium-dependent phospholipid binding  261285_at 1.0617 At3g05910 pectinacetylesterase, putative  253286_at 1.0555 At4g34260 catalytic  248596_at 1.0410 At5g49330 AtMYB111_AtMYB111 (myb domain protein 111); DNA binding / transcription  253140_at 1.0384 At4g35480 RHA3B_RHA3B (RING-H2 finger A3B); protein binding / zinc ion binding  RHL41_ZAT12RHL41 (RESPONSIVE TO HIGH LIGHT 41); nucleic acid binding / transcription factor/ zinc ion binding  MRH5_SHV3MRH5/SHV3 (morphogenesis of root hair 5);		+	At5g63130	
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ADT5_ADT5 (AROGENATE DEHYDRATASE 5); arogenate dehydratase/ prephenate dehydratase  264000_at	263799_at	1.0951	At2g24550	unknown [Populus trichocarpa x Populus deltoides] (GB:ABK96563.1)
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similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to unnamed protein product [Vitis vinifera] (GB:CAO69750.1)  250098_at				mitochondrial substrate carrier family protein
similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to 250098_at 1.0841 At5g17350 unknown [Populus trichocarpa] (GB:ABK95625.1) 262727_at 1.0669 At1g75800 pathogenesis-related thaumatin family protein ANNAT1_ATOXY5_OXY5_ANNAT1 (ANNEXIN ARABIDOPSIS 1); calcium ion 261285_at 1.0648 At1g35720 binding / calcium-dependent phospholipid binding 258750_at 1.0617 At3g05910 pectinacetylesterase, putative 253286_at 1.0555 At4g34260 catalytic 248596_at 1.0410 At5g49330 AtMYB111_AtMYB111 (myb domain protein 111); DNA binding / transcription 253140_at 1.0384 At4g35480 RHA3B_RHA3B (RING-H2 finger A3B); protein binding / zinc ion binding 247655_at 1.0351 At5g59820 binding / transcription factor/ zinc ion binding MRH5_SHV3_MRH5/SHV3 (morphogenesis of root hair 5);				similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G15358.1); similar to
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250098_at 1.0841 At5g17350 unknown [Populus trichocarpa] (GB:ABK95625.1)  262727_at 1.0669 At1g75800 pathogenesis-related thaumatin family protein  ANNAT1_ATOXY5_OXY5ANNAT1 (ANNEXIN ARABIDOPSIS 1); calcium ion  261285_at 1.0648 At1g35720 binding / calcium-dependent phospholipid binding  258750_at 1.0617 At3g05910 pectinacetylesterase, putative  253286_at 1.0555 At4g34260 catalytic  248596_at 1.0410 At5g49330 AtMYB111AtMYB111 (myb domain protein 111); DNA binding / transcription  253140_at 1.0384 At4g35480 RHA3BRHA3B (RING-H2 finger A3B); protein binding / zinc ion binding  247655_at 1.0351 At5g59820 binding / transcription factor/ zinc ion binding  MRH5_SHV3MRH5/SHV3 (morphogenesis of root hair 5);				similar to unknown protein [Arabidopsis thaliana] (TAIR:AT3G03280.1); similar to
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MRH5_SHV3MRH5/SHV3 (morphogenesis of root hair 5);	247655_at	1.0351	At5g59820	binding / transcription factor/ zinc ion binding
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	253925_at	1.0207	At4g26690	glycerophosphodiester phosphodiesterase/ kinase