

Table 2. 305_1hr_decrease

Probe set	Fold induction						Fold induction					
	Expression level (327,			Expression level (327,			quartz,			quartz,		
	1h)	305, 1h	295, 1h	1h)	1h)	305, 6h	295, 6h	6h	Identifier			
1 248676_at	595	0.3	0.1	0.1	614	1.2	1.1	0.3	AT5G48850	male sterility MS5 family		
2 247600_at	758	0.3	0.2	0.2	633	1.0	0.8	0.4	AT5G60890	myb transcription factor homolog (ATR1)		
3 252965_at	894	0.4	0.4	0.6	284	1.0	0.7	0.3	AT4G38860	auxin-induced (indole-3-acetic acid induced) protein, putative		
4 245668_at	372	0.4	0.6	1.2	292	0.9	0.9	0.8	AT1G28330	dormancy-associated protein -related		
5 266912_at	182	0.4	0.3	0.4	174	1.3	0.9	0.2	AT2G45900	expressed protein		
6 253317_at	439	0.5	0.4	0.4	823	1.1	0.8	0.3	AT4G33960	expressed protein		
7 266184_s_at	867	0.5	0.7	0.7	2061	1.0	0.9	1.2	AT2G38940	phosphate transporter (AtPT2)		
8 261895_at	1041	0.5	0.7	0.9	1210	1.0	0.7	0.5	AT1G80830	NRAMP metal ion transporter 1 (NRAMP1)		
9 259790_s_at	384	0.5	0.3	0.4	105	0.8	0.5	0.2	AT1G29430	auxin-induced (indole-3-acetic acid induced) protein family		
10 249008_at	956	0.6	0.4	0.5	725	0.9	0.7	0.4	AT5G44680	expressed protein		
11 251226_at	2161	0.6	0.7	0.5	1785	1.3	0.6	0.2	AT3G62680	proline-rich protein family		
12 252972_at	2190	0.6	0.4	0.5	1912	0.9	0.6	0.2	AT4G38840	auxin-induced (indole-3-acetic acid induced) protein, putative		
13 252833_at	723	0.6	0.7	0.4	548	1.3	0.6	0.2	AT4G40090	arabinogalactan-protein (AGP3)		
14 262844_at	639	0.6	0.6	0.8	544	1.1	1.0	0.7	AT1G14890	pectinesterase family		
15 267461_at	5382	0.6	0.6	0.8	1063	1.0	0.7	0.7	AT2G33830	auxin-regulated protein		
16 247377_at	890	0.6	0.6	1.0	2021	1.0	0.7	0.3	AT5G63180	polysaccharide lyase family 1 (pectate lyase)		
17 260668_at	589	0.6	0.5	0.6	117	0.6	0.9	0.7	AT1G19530	expressed protein		
18 245602_at	682	0.6	0.6	1.0	1002	1.1	0.9	0.7	AT4G14270	expressed protein		
19 248801_at	378	0.6	0.5	0.8	354	1.0	1.0	1.2	AT5G47370	homeobox-leucine zipper protein HAT2 (HD-ZIP protein 2)		
20 264348_at	2645	0.6	0.5	0.5	3724	0.9	0.7	0.5	AT1G12110	putative NPK1-related protein kinase 2		
21 264590_at	774	0.6	0.7	0.9	727	1.1	0.8	0.5	AT2G17710	expressed protein		
22 250054_at	510	0.6	1.0	1.0	530	1.2	1.5	3.5	AT5G17860	cation exchanger, putative (CAX7)		
23 251221_at	1922	0.6	1.1	2.3	5003	0.9	1.0	0.4	AT3G62550	expressed protein		
24 265471_at	3093	0.6	0.7	0.7	1068	1.1	0.5	0.6	AT2G37130	peroxidase, putative (ATP2a)		
25 260101_at	1695	0.7	0.8	0.6	651	1.1	1.0	0.9	AT1G73260	trypsin inhibitor -related		
26 246390_at	3415	0.7	0.6	0.5	2801	1.2	1.0	0.5	AT1G77330	1-aminocyclopropane-1-carboxylate oxidase (ACC oxidase), putative		
27 265444_s_at	5385	0.7	0.7	0.6	4280	1.2	1.1	1.1	AT2G37180	plasma membrane intrinsic protein 2C		
28 259541_at	917	0.7	0.6	0.7	757	1.1	1.0	0.5	AT1G20650	protein kinase -related		
29 255433_at	1536	0.7	0.6	0.6	1695	0.9	0.8	0.4	AT4G03210	xyloglucan endotransglycosylase, putative		

Fold induction for 305-1h is averaged from four independent ATH-1 array hybridizations, all others from two independent array hybridizations.

Annotation June 15, 2003.