

Table 2:

Probe set	Baseline expression level ^s	Fold-change	p-value	Annotation [#]
763.m00027_s_at ^a	0.01	28.97	0.024	hypothetical protein
120.m00098_x_at	0.01	14.35	0.011	hypothetical protein
329.m00054_at	0.01	13.06	0.017	hypothetical protein
253.m00081_x_at	0.02	10.61	0.041	hypothetical protein
16.m00301_at	0.01	9.75	0.006	hypothetical protein
228.m00060_x_at	0.02	9.43	0.007	AIG1 family protein, putative (GO:0004765)
244.m00069_x_at	0.01	9.23	0.021	hypothetical protein
371.m00031_x_at	0.01	8.69	0.006	BspA-like leucine rich repeat protein, putative
295.m00030_at	0.02	8.11	0.036	conserved hypothetical protein
64.m00173_x_at	0.01	7.95	0.043	BspA-like leucine rich repeat protein, putative
94.m00138_at	0.02	7.09	0.033	hypothetical protein
36.m00220_s_at ^b	0.02	5.78	0.005	hypothetical protein
432.m00029_x_at	0.08	5.32	0.001	conserved hypothetical protein
1087.m00005_at	0.03	5.12	0.038	hypothetical protein
401.m00029_at	1.11	4.97	0.022	protein kinase, putative (GO:0005524)
92.m00177_x_at	0.05	4.39	0.009	hypothetical protein
86.m00156_s_at ^c	0.09	4.06	0.036	glutamine cyclotransferase, putative
287.m00045_x_at	0.03	3.93	0.026	hypothetical protein
450.m00030_at	0.12	3.90	0.043	hypothetical protein
2.m00587_x_at	0.04	3.83	0.009	hypothetical protein
115.m00143_at*	0.12	3.64	0.001	transcription initiation factor TFIID, putative (GO:0003677)
420.m00021_x_at	0.03	3.59	0.007	hypothetical protein
27.m00245_at	0.03	3.55	0.017	hypothetical protein
103.m00185_at	6.55	3.32	0.023	Fe-hydrogenase, putative (GO:0005489)
194.m00103_at*	0.07	3.27	0.029	protein kinase, putative (GO:0005524)
26.m00274_x_at	0.07	3.22	0.039	hypothetical protein
935.m00010_at	0.11	3.15	0.009	hypothetical protein
93.m00158_s_at ^d	0.10	3.10	0.014	DNA mismatch repair protein mutS, putative (GO:0006298)
97.m00140_at*	0.08	3.10	0.006	hypothetical protein
138.m00086_x_at	4.75	3.09	0.023	hypothetical protein
66.m00163_s_at	0.03	3.08	0.010	hypothetical protein
12.m00286_at	0.04	3.06	0.025	hypothetical protein
295.m00044_x_at	0.07	3.04	0.017	conserved hypothetical protein
141.m00082_at*	0.32	2.98	0.010	protein kinase, putative (GO:0004672)
267.m00069_at	0.04	2.96	0.001	hypothetical protein
4.m00648_x_at	0.04	2.89	0.032	hypothetical protein
112.m00119_at	0.04	2.85	0.035	hypothetical protein
472.m00058_s_at ^e	0.08	2.83	0.008	hypothetical protein
204.m00098_x_at	0.22	2.77	0.008	hypothetical protein
583.m00011_at	0.41	2.69	0.026	hypothetical protein
62.m00174_x_at	0.06	2.66	0.037	hypothetical protein
55.m00164_at	0.07	2.63	0.002	hypothetical protein
273.m00075_x_at	0.05	2.59	0.042	hypothetical protein
249.m00077_x_at	0.06	2.55	0.049	hypothetical protein
289.m00092_at	0.05	2.53	0.014	hypothetical protein
464.m00030_at	0.05	2.53	0.010	hypothetical protein
1.m00665_at	1.11	2.53	0.046	hypothetical protein
51.m00150_x_at	0.05	2.49	0.009	hypothetical protein
159.m00103_x_at	0.43	2.44	0.044	protein kinase, putative (GO:0004672)
19.m00288_at	0.07	2.36	0.005	hypothetical protein
308.m00054_x_at	0.05	2.34	0.009	hypothetical protein
5.m00447_x_at	0.12	2.33	0.016	hypothetical protein
51.m00167_at	0.05	2.33	0.042	hypothetical protein
27.m00266_x_at	0.06	2.30	0.050	ELL complex EAP30 subunit, putative
222.m00072_at	0.09	2.30	0.040	SPRY domain protein
183.m00091_x_at	8.10	2.30	0.026	acyl-CoA synthetase, putative (GO:0008152)
289.m00067_s_at ^f	0.09	2.30	0.004	short chain dehydrogenase family protein
43.m00197_x_at	7.97	2.24	0.024	cell adhesion protein (GO:0007155)
54.m00234_s_at ^g	1.09	2.22	0.037	hypothetical protein (GO:0005554)
4.m00613_x_at	0.12	2.21	0.007	hypothetical protein
48.m00224_s_at ^h	0.71	2.21	0.018	acetyltransferase, putative
75.m00156_x_at	0.11	2.17	0.007	myotubularin, putative
25.m00258_x_at	0.38	2.16	0.032	hypothetical protein
34.m00243_x_at	4.14	2.15	0.028	glucosamine 6-phosphate N-acetyltransferase, putative (GO:0008080)
314.m00052_x_at	0.07	2.12	0.041	hypothetical protein
21.m00282_at	0.21	2.08	0.038	hypothetical protein
528.m00022_at	0.10	2.03	0.018	hypothetical protein
48.m00192_at	0.17	2.01	0.013	hypothetical protein