

AGI code	Name	Log2FC	pvalue	Description
AT1G27570		3.55276	1.1E-06	phosphatidylinositol 3- and 4-kinase family protein
AT4G31650		1.9492	0.00851	transcriptional factor B3 family protein
AT4G29200	DEG3	1.89793	6.3E-25	over-expressed by salt stress
AT2G46420		1.84373	6.6E-60	helicase with zinc finger protein
AT1G17580	MYA1	1.62722	3.7E-14	member of the type XI myosin protein family
AT3G58640		1.56124	1.1E-24	MAPKKK-like protein
AT1G59890	SNL5	1.53126	1.8E-08	related to histone deacetylation
AT2G19390		1.51376	3.7E-10	serine/arginine repetitive matrix protein
AT1G47500	RBP47C	1.49849	2.2E-24	RNA-binding protein
AT1G49600	RBP47A	1.4612	8E-170	RNA-binding protein
AT5G42670		1.45621	0.00365	Agnet domain-containing protein
AT4G10730		1.39714	4.7E-16	protein kinase superfamily protein
AT3G51120		1.36742	1.4E-21	zinc finger CCCH domain-containing protein
AT3G12200	NEK7	1.24968	1.1E-09	member of the NIMA-related serine/threonine kinases
AT5G24430		1.24295	5.2E-20	calcium-dependent protein kinase (CDPK) family protein
AT1G07705	NOT2A	1.2257	0.00019	a negative regulator of translation
AT5G20320	DCL4	1.22428	6.2E-05	related to RNA processing
AT5G43810	AGO10	1.22149	0.00199	related to RNA processing
AT1G08600	ATRX	1.21933	3.8E-16	related to chromatin remodeling
AT3G24520	HSFC1	1.19593	6.4E-47	HSF type transcription factor related to seed dormancy
AT1G47490	RBP47C	1.18058	1.7E-24	RNA-binding protein
AT5G50350		1.16848	1.1E-09	hypothetical protein
AT3G50780	BTBPOZ	1.14854	0.00013	BTB/POZ domain protein related to protein ubiquitination
AT3G24650	ABI3	1.07477	7.2E-82	B3 type transcription factor related to seed dormancy
AT4G11330	MPK5	1.07155	0.00206	MAP kinase
AT5G38600		1.00075	3.2E-07	proline-rich spliceosome-associated (PSP) family protein