

gene	gene.name	#	baseMean	#	log2FoldChange	#	lfcSE	#	pvalue	#	padj	biological function
CAALFM_C102590CA	pyridoxine biosynthesis protein		28052.795		2.547754759		0.2527925177		3.85E-25		2.57E-22	amino acid metabolic process, pyridoxal phosphate
CAALFM_C102600WA	putative pyridoxal 5'-phosphate synthase		2442.209911		1.189647103		0.2844172839		1.16E-06		0.0005787418265	glutamine metabolic process, pyridoxal phosphate
CAALFM_C203370WA	trifunctional hydroxymethylpyrimidine kinase/phos		3682.204282		4.446556717		0.2283812204		1.12E-85		1.12E-82	phosphorylation, thiamine biosynthetic process
CAALFM_C204580WA	bifunctional (2E,6E)-farnesyl diphosphate synthase		5087.68292		1.013875747		0.1744538363		2.80E-10		1.53E-07	ergosterol biosynthetic process, farnesyl diphosph
CAALFM_C302860WA	bifunctional hydroxyethylthiazole kinase/thiamine- π		4665.24953		4.681733708		0.1999233449		1.68E-122		3.36E-119	phosphorylation, thiamine biosynthetic process, thi
CAALFM_C302870CA	uncharacterized protein		303.6545513		1.209277054		0.3912773681		6.79E-05		0.03136513369	amino acid transmembrane transport, cell-abiotic s
CAALFM_C305130CA	thiamine thiazole synthase		148044.5651		11.54563842		0.5179486034		2.79E-111		4.19E-108	thiamine biosynthetic process, thiazole biosynthetic
CAALFM_C404220WA	Duo1p		320.1559081		1.710978788		0.2576203805		1.40E-12		8.41E-10	filamentous growth, mitotic spindle organization
CAALFM_C404230WA	uncharacterized protein		7902.060114		5.162791222		0.2044699808		3.91E-142		2.35E-138	transmembrane transporter activity
CAALFM_C503480CA	uncharacterized protein		19328.82792		3.887612755		0.1990817987		1.90E-86		2.29E-83	cellular response to metal ion, cellular response to
CAALFM_CR09290WA	4-amino-5-hydroxymethyl-2-methylpyrimidine phos		71545.15854		9.309949857		0.4968035323		1.50E-79		1.29E-76	thiamine biosynthetic process, thiamine diphospha
CAALFM_CR09350CA	uncharacterized protein		1174.742033		8.173936016		0.3380966931		4.21E-129		1.26E-125	thiamine metabolic process
CAALFM_CR09360WA	Fcy24p		7084.390493		6.345570528		0.4999566475		3.51E-38		2.63E-35	transmembrane transporter activity