

coding region	MOWSE score	name	description	GenDB ID
cg0040	123 72 80 88 89 90		PUTATIVE SECRETED PROTEIN	2631
cg0482	74	gpmA	PHOSPHOGLYCEROMUTASE 1 (EC 5.4.2.1)	127
cg0700	73	guaB3	IMP dehydrogenase / GMP reductase C terminus	3013
cg0755	154	metY	O-acetylhomoserine sulfhydrylase	1136
cg1111	76 77 79 80	eno	ENOLASE (EC 4.2.1.11) (2-PHOSPHOGLYCERATE DEHYDRAT	973
cg1145	85	fum	FUMARATE HYDRATASE (EC 4.2.1.2)	5
cg1380	72	ssuA	ABC-type aliphatic sulfonate transporter, substrate-binding protein	1087
cg1514	70		secreted protein	1513
cg1735	121 130 141 166 170 173		secreted cell wall-associated hydrolase (invasion-associated protein)	2421
cg1763	101 75	sufD	components of an uncharacterized iron-regulated ABC-type transporter	852
cg1774	69	tkt	TRANSKETOLASE (EC 2.2.1.1)	2589
cg1790	67 92	pgk	PHOSPHOGLYCERATE KINASE (EC 2.7.2.3)	489
cg1791	104 111	gap	GLYCERALDEHYDE-3-PHOSPHATE DEHYDROGENASE (EC 1.2.1)	1677

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cg2052	66 74 75 77		putative secreted protein	524
cg2057	68		putative secreted protein	376
cg2221	111	tsf	TRANSLATION ELONGATION FACTOR TS (EF-TS)	2983
cg2361	75 75		Cell division initiation protein - Antigen 84 homolog	2846
cg2401	88		SECRETED PROTEIN NLP/P60 FAMILY, PUTATIVE PEPTIDOGLYCAN LYTIC PROTEIN	4414
cg2591	77	dkgA	2,5-DIKETO-D-GLUCONIC ACID REDUCTASE (EC 1.1.1.-)	2755
cg2833	122 136 154 75 81 94 98	cysK	O-Acetylserine (Thiol)-Lyase	532
cg2840	80 82	actA	BUTYRYL-COA:ACETATE COENZYME A TRANSFERASE	2187
cg3049	98	fpr1	NADPH-dependent ferredoxin reductase	277
cg3096	114 83		ALDEHYDE DEHYDROGENASE (EC 1.2.1.3)	4002
cg3100	121 127 141 72 76	dnaK	Heat shock protein hsp70	2581
cg3115	99	cysD	sulfate adenylyltransferase subunit 2	682
cg3119	72	fpr2	NADPH-dependent ferredoxin reductase	2569
cg3182	69 83	cop1	Trehalose corynomycolyl transferase2	3095