

GENE	GO BIOLOGICAL PROCESS
CG32816	biological_process no biological data available
CG13369	D-ribose metabolic process inferred from electronic annotation with InterPro:IPR002139
CG13362	-
CG14635	-
CG14633	-
ThiA-5-2	transcription initiation from RNA polymerase II promoter inferred from sequence or structural similarity with HGNC_gene:GTF2A2; HGNC:4647; OMIM:600519 AND inferred from sequence or structural similarity with SGD_LOCUS:TOA2; SGD:S0001541
CG14631	regulation of transcription from RNA polymerase II promoter non-traceable author statement
CG33082	-
CG11381	-
CG14781	mitotic spindle elongation inferred from mutant phenotype
CG32810	ion transport inferred from sequence or structural similarity with NCBI_NP:508239
CG32803	potassium ion transport inferred from electronic annotation with InterPro:IPR003131
CG3857	-
CG4325	-
CG14054	-
CG4061	RNA processing inferred from electronic annotation with InterPro:IPR013791, InterPro:IPR017770
CG3391	transport inferred from electronic annotation with InterPro:IPR01071
CG2882	-
CG10793	response to oxidative stress inferred from electronic annotation with InterPro:IPR002016
CG14269	-
CG16782	-
Mnt	regulation of transcription inferred from sequence or structural similarity with dm
	phagocytosis_enquilmnt inferred from mutant phenotype
CG15671	-
CG15676	-
CG15677	-
CG15786	-
CG15768	proteolysis inferred from electronic annotation with InterPro:IPR000209, InterPro:IPR000834
CG15764	-
CG12236	-
CG8041	proteolysis inferred from electronic annotation with InterPro:IPR001254, InterPro:IPR001314, InterPro:IPR018114
CG17171	-
CG3226	-
CG4575	regulation of transcription_DNA-dependent inferred from electronic annotation with InterPro:IPR004827, InterPro:IPR011700
inx7	-
CG1999	-
CG15330	-
CG2256	-
CG2147	-
CG12661	-
CG10970	-
Dipt	-
CG12057	-
CG34339	-
CG15317	-
Gaa	post-Golgi vesicle-mediated transport inferred from sequence or structural similarity with UniProtKB_Q9UJY5, UniProt_Q9UJY4, UniProt_Q9NZ52
	intracellular protein transport inferred from electronic annotation with InterPro:IPR002014, InterPro:IPR004152, InterPro:IPR008152, InterPro:IPR013041
CG1791	signal transduction inferred from electronic annotation with InterPro:IPR002181, InterPro:IPR014716
Slp	-
l11G0289	-
CG2888	-
RaxX2	regulation of transcription_DNA-dependent inferred from electronic annotation with InterPro:IPR002078
	small GTPase mediated signal transduction inferred from electronic annotation with InterPro:IPR001806, InterPro:IPR003579
	protein transport inferred from electronic annotation with InterPro:IPR003579
CG15296	-
CG2124	-
CG12637	-
CG15209	-
CG17333	pentose-phosphate shunt non-traceable author statement
CG11106	-
CG11756	-
CG1738	-
CG9360	metabolic process inferred from electronic annotation with InterPro:IPR002198, InterPro:IPR002347, InterPro:IPR016040
CG15739	metabolic process inferred from electronic annotation with InterPro:IPR005834, InterPro:IPR006357
CG11366	-
CG15925	-
CG1463	-
CG2543	nitrohydroxyglycylutamate biosynthetic process inferred from sequence or structural similarity with UniProtKB_Q05932
mRpl49	translational initiation inferred from electronic annotation with InterPro:IPR001950
CG32643	-
CG4351	-
CG15747	-
CG12725	protein modification process inferred from electronic annotation with InterPro:IPR000626
CG12481	-
CG13403	-
CG12540	-
nmdyn-D6	GTP biosynthetic process inferred from electronic annotation with InterPro:IPR001564
	UTP biosynthetic process inferred from electronic annotation with InterPro:IPR001564
	GTP biosynthetic process inferred from electronic annotation with InterPro:IPR001564
CG15641	transport inferred from electronic annotation with InterPro:IPR001588
CG15801	-
CG12395	-
CG34015	biological_process no biological data available
CG4653	proteolysis inferred from electronic annotation with InterPro:IPR001254, InterPro:IPR001314, InterPro:IPR018114
CG4789	regulation of transcription_DNA-dependent inferred from electronic annotation with InterPro:IPR002078
CG13004	-
CG4955	-
CG18258	lipid metabolic process inferred from electronic annotation with InterPro:IPR000734
CG8915	-
CG8408	biological_process no biological data available
CG12672	-
Her	negative regulation of transcription from RNA polymerase II promoter inferred from sequence or structural similarity with MGD_Hes1; MGI:MGI:104853
CG15043	-
CG15047	-
CG6873	-
CG34329	-
CG1504	-
Obp19a	olfactory behavior inferred from sequence or structural similarity with UniProtKB_Q02372
	response to pheromone inferred from sequence or structural similarity with UniProtKB_Q02372
	sensory perception of chemical stimulus inferred from sequence or structural similarity with UniProtKB_P34174
	transport inferred from electronic annotation with InterPro:IPR006170
CG1324	-
CG1304	proteolysis non-traceable author statement
CG1722	-
CG41106	biological_process no biological data available
FucTC	protein amino acid glycosylation inferred from electronic annotation with InterPro:IPR001503
Rab21	small GTPase mediated signal transduction inferred from electronic annotation with InterPro:IPR001806, InterPro:IPR003579
	protein transport inferred from electronic annotation with InterPro:IPR003579
CG13865	biological_process no biological data available