

GENE	GO BIOLOGICAL PROCESS
CG31516	biological process no biological data available
CG14639	.
CG41461	.
CG41468	.
CG31525	biological process no biological data available
CG31526	biological process no biological data available
CG41516	.
CG14659	.
CG14662	.
CG12590	.
CG12161	ubiquitin-dependent protein catabolic process inferred from electronic annotation with InterPro:IPR000243, InterPro:IPR001353
mia	transcription initiation from RNA polymerase II promoter inferred from sequence or structural similarity with HGNC_gene:TAF6L; HGNC:17305; OMIM:602946 AND inferred from sequence or structural similarity with SGD_LOCUS:TAF60; SGD:S0003080 male meiosis I inferred from mutant phenotype spermatogenesis inferred from mutant phenotype meiosis I traceable author statement spermatid development traceable author statement regulation of transcription factor activity inferred from electronic annotation with InterPro:IPR011442
CG31554	biological process no biological data available
CG1075	proteolysis inferred from electronic annotation with InterPro:IPR000668
CG15185	.
CG1138	.
nx14	RNA export from nucleus inferred from sequence or structural similarity with NCBI_NP:071336
Mst84Dd	.
CG11977	.
CG33630	biological process no biological data available
CG34108	biological process no biological data available
CG14694	transport inferred from electronic annotation with InterPro:IPR002666
CG14708	.
ht	negative regulation of protein amino acid dephosphorylation inferred from direct assay
CG10013	.
CG10096	metabolic process inferred from electronic annotation with InterPro:IPR016040
CG17207	biological process no biological data available
CG33929	biological process no biological data available
CG10909	RNA processing inferred from sequence or structural similarity with SGD_LOCUS:NOP1; SGD:S0002172
CG14394	.
CG14370	.
CG14369	.
CG9929	.
CG14357	.
CG9616	.
CG14851	.
FK506-bp1	protein folding inferred from sequence or structural similarity with UniProtKB:Q26486
CG5399	.
CG34274	.
CG12784	.
CG5614	.
ird5	positive regulation of antibacterial peptide biosynthetic process inferred from mutant phenotype protein catabolic process inferred from direct assay protein amino acid phosphorylation non-traceable author statement Toll signaling pathway traceable author statement dorsal/ventral axis specification non-traceable author statement antibacterial humoral response inferred from mutant phenotype antimicrobial humoral response non-traceable author statement protein folding inferred from electronic annotation with InterPro:IPR004127
CG31287	protein folding inferred from electronic annotation with InterPro:IPR004127
Mst89B	.
CG17930	transport inferred from electronic annotation with InterPro:IPR005828
CG10317	.
Actn3	actin filament bundle formation inferred from sequence or structural similarity with EMBL:AF093775
CG5866	.
Sns5	pupal adhesion traceable author statement
CG34281	.
CG14315	.
CG14313	.
CG7669	.
CG14306	.
CG4538	ATP-dependent proteolysis inferred from sequence or structural similarity with UniProtKB:O76031 protein transport inferred from electronic annotation with InterPro:IPR004487
Pros28.1A	ubiquitin-dependent protein catabolic process inferred from electronic annotation with InterPro:IPR000426, InterPro:IPR001353
TotC	response to stress inferred from sequence or structural similarity with TotA
CheB93b	.
CG6690	oxidation reduction inferred from electronic annotation with InterPro:IPR006863 cell redox homeostasis inferred from electronic annotation with InterPro:IPR013766, InterPro:IPR017936
BG4	defense response non-traceable author statement signal transduction non-traceable author statement immune response inferred from mutant phenotype apoptosis non-traceable author statement sperm individualization inferred from mutant phenotype defense response to Gram-negative bacterium traceable author statement positive regulation of antibacterial peptide biosynthetic process traceable author statement
CG17625	.
CG31281	.
p53	DNA damage response, signal transduction by p53 class mediator traceable author statement DNA damage response, signal transduction resulting in induction of apoptosis traceable author statement apoptosis traceable author statement cell death inferred from mutant phenotype response to radiation inferred from mutant phenotype DNA damage response, signal transduction by p53 class mediator resulting in induction of apoptosis inferred from mutant phenotype response to DNA damage stimulus inferred from mutant phenotype regulation of transcription from RNA polymerase II promoter traceable author statement determination of adult life span inferred from mutant phenotype issue regeneration inferred from mutant phenotype response to UV inferred from mutant phenotype caspase activation inferred from mutant phenotype response to ionizing radiation inferred from mutant phenotype positive regulation of gene-specific transcription inferred from mutant phenotype positive regulation of cell cycle inferred from genetic interaction with CoVa
CG31457	.
CG31515	.
CG10177	protein amino acid phosphorylation non-traceable author statement intracellular signaling cascade inferred from electronic annotation with InterPro:IPR003533
CG10164	.
CG16710	proteolysis non-traceable author statement
CG18754	proteolysis non-traceable author statement
CG10254	regulation of protein metabolic process inferred from electronic annotation with InterPro:IPR000608 post-translational protein modification inferred from electronic annotation with InterPro:IPR000608
gav	telomere capping inferred from expression pattern telomere maintenance inferred from mutant phenotype
CG31128	biological process no biological data available
bam	gamete generation inferred from mutant phenotype

	cystoblast division inferred from mutant phenotype
	fusome organization inferred from mutant phenotype
	spermatogenesis traceable author statement
	germ-line stem cell division inferred from mutant phenotype
	germanium-derived female germ-line cyst formation inferred from mutant phenotype
	cell fate specification traceable author statement
	germ cell development traceable author statement
	male meiosis traceable author statement
	female germ-line cyst formation traceable author statement
	cell fate determination traceable author statement
	spectrosome organization traceable author statement
	spermatogonial cell division traceable author statement
	vesicle targeting to fusome traceable author statement
	oogenesis inferred from mutant phenotype
CG11889	-
BG642167	-
Mat57Db	oviposition non-traceable author statement
	post-mating behavior non-traceable author statement
CG14545	-
CG34130	biological process no biological data available
CG14244	-
CG14262	-
CG5402	-
CG12879	-
Acp98AB	negative regulation of female receptivity, post-mating non-traceable author statement
CG33213	biological process no biological data available
CG5527	proteolysis inferred from electronic annotation with InterPro:IPR000718, InterPro:IPR008753, InterPro:IPR018497
CG34436	-
CG15510	-
Jon99Fi	proteolysis inferred from electronic annotation with InterPro:IPR001254, InterPro:IPR001314, InterPro:IPR018114
CG2267	-
CG31013	protein metabolic process inferred from electronic annotation with InterPro:IPR006620
	oxidation reduction inferred from electronic annotation with InterPro:IPR013547
CG1340	translational initiation inferred from sequence or structural similarity
CG31008	-
Prosalpha3T	ubiquitin-dependent protein catabolic process inferred from electronic annotation with InterPro:IPR000426, InterPro:IPR001353
CG18672	one-carbon compound metabolic process inferred from electronic annotation with InterPro:IPR018338
CG1792	-