

ESTMD

EST Model Database

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GO Sequence Search Result

SequenceID	GO_ID	GO_Term	Type	HitID	Gene_Symbol
Contig1	GO:0003735	structural constituent of ribosome	molecular_function	FBgn0004413	RpS26
Contig1	GO:0006412	protein biosynthesis	biological_process	FBgn0004413	RpS26
Contig1	GO:0005843	cytosolic small ribosomal subunit (sensu Eukarya)	cellular_component	FBgn0004413	RpS26
Contig1	GO:0005840	ribosome	cellular_component	FBgn0004413	RpS26
Contig10	GO:0006412	protein biosynthesis	biological_process	FBgn0011272	RpL13
Contig10	GO:0005842	cytosolic large ribosomal subunit (sensu Eukarya)	cellular_component	FBgn0011272	RpL13
Contig10	GO:0005840	ribosome	cellular_component	FBgn0011272	RpL13
Contig10	GO:0003735	structural constituent of ribosome	molecular_function	FBgn0011272	RpL13
Contig106	GO:0004772	sterol O-acyltransferase	molecular_function	FBgn0031327	CG5397
Contig35	GO:0006096	glycolysis	biological_process	FBgn0001258	ImpL3
Contig35	GO:0004459	L-lactate dehydrogenase	molecular_function	FBgn0001258	ImpL3