

**\_F\_TULARENSIS\_LVS\_NEUTROPHIL\_DN, GSE37416\_12H\_VS\_24H\_F\_TULARENSIS\_LVS\_NEUTROPHIL\_DN**

GOCC\_AZUROPHIL\_GRANULE, GOCC\_AZUROPHIL\_GRANULE  
GOBP\_VACUOLAR\_TRANSPORT, GOBP\_VACUOLAR\_TRANSPORT  
MIR6739\_3P, MIR6739\_3P  
GOBP\_LYSOSOMAL\_TRANSPORT, GOBP\_LYSOSOMAL\_TRANSPORT  
GOBP\_UBIQUITIN\_DEPENDENT\_ERAD\_PATHWAY, GOBP\_UBIQUITIN\_DEPENDENT\_ERAD\_PATHWAY  
HP\_GENU\_VALGUM, HP\_GENU\_VALGUM  
FAELT\_B\_CLL\_WITH\_VH\_REARRANGEMENTS\_UP, FAELT\_B\_CLL\_WITH\_VH\_REARRANGEMENTS\_UP  
HP\_EXCESSIVE\_SALIVATION, HP\_EXCESSIVE\_SALIVATION  
GOCC\_PRESYNAPTIC\_ACTIVE\_ZONE, GOCC\_PRESYNAPTIC\_ACTIVE\_ZONE  
GOBP\_NEGATIVE\_REGULATION\_OF\_LEUKOCYTE\_MIGRATION, GOBP\_NEGATIVE\_REGULATION\_OF\_  
HP\_ANONYCHIA, HP\_ANONYCHIA  
MIR23A\_5P, MIR23A\_5P  
GOBP\_MONOAMINE\_TRANSPORT, GOBP\_MONOAMINE\_TRANSPORT  
MODULE\_418, MODULE\_418  
HP\_SHORT\_UPPER\_LIP, HP\_SHORT\_UPPER\_LIP  
MODULE\_455, MODULE\_455  
HP\_RENOVASCULAR\_HYPERTENSION, HP\_RENOVASCULAR\_HYPERTENSION  
HP\_DELAYED\_FINE\_MOTOR\_DEVELOPMENT, HP\_DELAYED\_FINE\_MOTOR\_DEVELOPMENT  
KEGG\_GLYCOSPHINGOLIPID\_BIOSYNTHESIS\_GLOBO\_SERIES, KEGG\_GLYCOSPHINGOLIPID\_BIOSYNT  
MYF6\_TARGET\_GENES, MYF6\_TARGET\_GENES