tissue_15_pos_Neutrophil_tissue−lungvsmln_bu
translation at postsynapse
cytokinesis
regulation of lamellipodium organization
regulation of amyloid precursor protein catabolic process
regulation of regulated secretory pathway
miRNA metabolic process
regulation of potassium ion transmembrane transport
negative regulation of amide metabolic process
ubiquitin−dependent protein catabolic process via the multivesicular body sorting pathway
regulation of receptor localization to synapse
modification of postsynaptic actin cytoskeleton
positive regulation of blood pressure
membrane docking
synapse maturation
protein localization to cell−cell junction
pallium development
positive regulation of mRNA cis splicing, via spliceosome
mast cell secretory granule organization
positive regulation of protein K48−linked ubiquitination
negative regulation of hydrolase activity
response to leptin
Arp2/3 complex−mediated actin nucleation
motor neuron apoptotic process
interleukin−11−mediated signaling pathway
negative regulation of stem cell differentiation
negative regulation of vascular associated smooth muscle cell migration
organophosphate metabolic process
positive regulation of T cell activation
type II interferon production
negative regulation of B cell activation
antigen processing and presentation of peptide or polysaccharide antigen via MHC class II
peptide antigen assembly with MHC class II protein complex
regulation of JNK cascade
heparin proteoglycan metabolic process
sebaceous gland development
positive regulation of growth rate
isoprenoid metabolic process
small molecule metabolic process
mitochondrial translation
thioester metabolic process
acute inflammatory response
glomerulus development
positive regulation of mononuclear cell proliferation
GO term
−2
−1
0
1
2
3
NES
−log10(p)
2
3
4
5
6
NES
2
1
0
−1
−2