tissue_28_neg_iMO1_tissue−bloodvsmln_bubble
glutathione metabolic process
sulfur compound metabolic process
olefinic compound metabolic process
regulation of bone mineralization
xenobiotic metabolic process
fatty acid oxidation
organic acid biosynthetic process
actomyosin contractile ring organization
amide metabolic process
monocyte activation
venous blood vessel development
NMDA selective glutamate receptor signaling pathway
respiratory basal cell differentiation
parturition
myelin maintenance
glycoprotein metabolic process
observational learning
neuron intrinsic apoptotic signaling pathway in response to endoplasmic reticulum stress
endothelin receptor signaling pathway
positive regulation of ATP−dependent activity
glycerol catabolic process
negative regulation of leukocyte migration
modification of postsynaptic structure
negative regulation of potassium ion transport
establishment of mitotic spindle orientation
regulation of cAMP−dependent protein kinase activity
regulation of mRNA processing
positive regulation of heat generation
regulation of type I interferon production
endodermal digestive tract morphogenesis
amygdala development
RNA transport
regulation of chromatin organization
cell killing
nuclear−transcribed mRNA catabolic process
positive regulation of humoral immune response
regulation of defense response to virus by host
cellular response to interferon−beta
antigen processing and presentation of endogenous antigen
protein refolding
peptide antigen assembly with MHC protein complex
RNA localization
GO term
−3
−2
−1
0
1
2
NES
−log10(p)
3
4
5
6
NES
2
1
0
−1
−2