mtb_mln_28_Neutrophil_Mtb−posvsneg_
modified amino acid metabolic process
response to pH
aerobic respiration
aldehyde metabolic process
carbohydrate derivative metabolic process
mitochondrial transport
oxoacid metabolic process
binding of sperm to zona pellucida
purine−containing compound biosynthetic process
negative regulation of peptidase activity
amide metabolic process
regulation of chemokine (C−X−C motif) ligand 2 production
terpenoid metabolic process
nerve development
regulation of coagulation
nucleoside phosphate biosynthetic process
positive regulation of establishment of protein localization to telomere
sensory perception of pain
sulfur compound biosynthetic process
amino acid metabolic process
monosaccharide metabolic process
locomotory behavior
lactone metabolic process
semaphorin−plexin signaling pathway
regulation of plasma lipoprotein particle levels
cysteine transmembrane transport
retrograde trans−synaptic signaling
protein maturation
cellular response to cAMP
mRNA splice site recognition
dichotomous subdivision of terminal units involved in salivary gland branching
negative regulation of non−canonical NF−kappaB signal transduction
regulation of Fc receptor mediated stimulatory signaling pathway
regulation of actin filament length
respiratory burst involved in defense response
response to nerve growth factor
interleukin−4−mediated signaling pathway
platelet degranulation
membrane raft localization
positive regulation of nuclear−transcribed mRNA catabolic process, deadenylation−dependent decay
modification of postsynaptic actin cytoskeleton
early endosome to recycling endosome transport
GO term
−2
−1
0
1
2
NES
−log10(p)
4
6
8
NES
2
1
0
−1
−2