mtb\_mln\_15\_iMO1\_Mtb-posvsneg\_bubble.pdf mitochondrial gene expression acute-phase response oxidative phosphorylation organic acid transport mitochondrial respiratory chain complex assembly heat generation ribonucleotide biosynthetic process prostanoid biosynthetic process response to fatty acid modified amino acid transport myotube differentiation involved in skeletal muscle regeneration peptide antigen assembly with MHC class II protein complex positive regulation of smooth muscle contraction -log10(p) stem cell fate commitment protein kinase C deactivation 2.5 3.0 glomerular endothelium development 3.5 regulation of tolerance induction to self antigen dendritic transport chorio-allantoic fusion GO term positive regulation of cellular response to transforming growth factor beta stimulus 5.0 sodium ion export across plasma membrane positive regulation of lamellipodium organization **NES** Golgi reassembly macrophage migration 0 regulation of long-term neuronal synaptic plasticity negative regulation of DNA binding cytoplasmic pattern recognition receptor signaling pathway macrophage differentiation regulation of actin filament length mature B cell differentiation involved in immune response type II interferon-mediated signaling pathway positive regulation of tissue remodeling regulation of interleukin-23 production N-terminal peptidyl-lysine acetylation alternative mRNA splicing, via spliceosome defense response to virus translation at postsynapse negative regulation of viral genome replication establishment or maintenance of epithelial cell apical/basal polarity receptor metabolic process 2 -3 -2 0 **NES**