Macrophages_LYVE1 - IMM - Colon - ActiveCD_vs_Control protein phosphorylation - GO:0006468 - p: 1.69e-04 *** phosphorylation - GO:0016310 - p: 9.76e-05 *** negative regulation of cell adhesion - GO:0007162 - p: 2.30e-04 *** alpha-beta T cell activation - GO:0046631 - p: 1.73e-04 *** lymphocyte activation involved in immune response - GO:0002285 - p: 1.44e-04 *** alpha-beta T cell differentiation - GO:0046632 - p: 2.35e-04 *** CD4-positive, alpha-beta T cell activation - GO:0035710 - p: 5.58e-05 *** generation of neurons - GO:0048699 - p: 2.07e-04 *** response to cytokine - GO:0034097 - p: 1.53e-04 *** positive regulation of intracellular signal transd... - GO:1902533 - p: 8.40e-05 *** cellular response to endogenous stimulus - GO:0071495 - p: 1.55e-04 *** positive regulation of developmental process - GO:0051094 - p: 8.86e-05 *** system process - GO:0003008 - p: 5.70e-05 *** response to nitrogen compound - GO:1901698 - p: 3.06e-05 *** response to hormone - GO:0009725 - p: 6.49e-05 *** Cluster regulation of immune response - GO:0050776 - p: 2.72e-05 *** hemopoiesis - GO:0030097 - p: 1.80e-05 *** anatomical structure formation involved in morphog... - GO:0048646 - p: 6.26e-05 *** neuron development - GO:0048666 - p: 1.35e-04 *** GO Term regulation of defense response - GO:0031347 - p: 4.07e-05 *** innate immune response - GO:0045087 - p: 3.63e-06 *** Gene Count regulation of response to external stimulus - GO:0032101 - p: 1.87e-06 *** 100 regulation of response to biotic stimulus - GO:0002831 - p: 2.64e-05 *** response to peptide hormone - GO:0043434 - p: 2.39e-04 ** perinuclear region of cytoplasm - GO:0048471 - p: 8.86e-05 *** sequence-specific DNA binding - GO:0043565 - p: 6.15e-05 *** RNA polymerase II transcription regulatory region ... – GO:0000977 – p: 1.87e–04 *** DNA-binding transcription factor activity, RNA pol... - GO:0000981 - p: 4.03e-04 *** DNA-binding transcription factor activity - GO:0003700 - p: 2.21e-04 *** double-stranded DNA binding - GO:0003690 - p: 1.10e-05 *** transcription regulatory region nucleic acid bindi... - GO:0001067 - p: 2.16e-05 *** transcription cis-regulatory region binding - GO:0000976 - p: 2.16e-05 *** sequence-specific double-stranded DNA binding - GO:1990837 - p: 7.43e-06 *** protein kinase activity - GO:0004672 - p: 1.76e-04 *** phosphotransferase activity, alcohol group as acce... - GO:0016773 - p: 3.06e-05 *** histone kinase activity - GO:0035173 - p: 1.71e-04 *** histone H3 kinase activity - GO:0140996 - p: 1.69e-04 *** histone H2AX kinase activity - GO:0141003 - p: 1.33e-04 *** histone modifying activity - GO:0140993 - p: 5.58e-07 *** 1.4 1.5 1.3 1.6 Absolute NES