Stem_cells_OLFM4_PCNA - EPI - Colon - InactiveCD_vs_Control cellular response to hormone stimulus - GO:0032870 - p: 3.20e-07 *** tube morphogenesis - GO:0035239 - p: 3.37e-07 *** positive regulation of cell differentiation - GO:0045597 - p: 2.18e-07 *** regulation of phosphorus metabolic process - GO:0051174 - p: 2.84e-08 *** regulation of phosphate metabolic process - GO:0019220 - p: 2.74e-08 *** regulation of anatomical structure morphogenesis - GO:0022603 - p: 1.02e-07 *** blood vessel development - GO:0001568 - p: 2.56e-07 *** regulation of phosphorylation - GO:0042325 - p: 3.63e-08 *** negative regulation of developmental process - GO:0051093 - p: 6.20e-09 *** regulation of protein phosphorylation - GO:0001932 - p: 3.73e-08 *** epithelial cell differentiation - GO:0030855 - p: 9.11e-09 *** regulation of cell development - GO:0060284 - p: 4.71e-10 *** negative regulation of cell differentiation - GO:0045596 - p: 2.32e-09 *** lymphocyte activation - GO:0046649 - p: 1.25e-07 *** BP leukocyte activation - GO:0045321 - p: 1.70e-08 *** Cluster cell-cell adhesion - GO:0098609 - p: 2.70e-10 *** regulation of cell adhesion - GO:0030155 - p: 1.00e-10 *** regulation of lymphocyte activation - GO:0051249 - p: 4.30e-08 *** regulation of cell activation - GO:0050865 - p: 1.09e-09 *** regulation of leukocyte activation - GO:0002694 - p: 2.96e-09 *** positive regulation of cell adhesion - GO:0045785 - p: 2.77e-09 *** regulation of T cell activation - GO:0050863 - p: 4.31e-08 *** regulation of cell-cell adhesion - GO:0022407 - p: 9.47e-10 *** Gene Count leukocyte cell-cell adhesion - GO:0007159 - p: 7.69e-09 *** regulation of leukocyte cell-cell adhesion - GO:1903037 - p: 3.37e-08 *** positive regulation of leukocyte cell-cell adhesio... - GO:1903039 - p: 1.80e-07 *** positive regulation of cell-cell adhesion - GO:0022409 - p: 2.14e-08 *** 0 1 adaptive immune response - GO:0002250 - p: 2.86e-09 *** transcription regulator complex - GO:0005667 - p: 1.15e-03 ** perinuclear region of cytoplasm - GO:0048471 - p: 4.44e-04 *** CC somatodendritic compartment - GO:0036477 - p: 1.43e-03 ** axon - GO:0030424 - p: 7.16e-05 *** side of membrane - GO:0098552 - p: 1.05e-04 *** phosphotransferase activity, alcohol group as acce... - GO:0016773 - p: 1.58e-05 *** histone modifying activity - GO:0140993 - p: 1.16e-05 *** protein kinase activity - GO:0004672 - p: 2.25e-06 *** histone H2AX kinase activity - GO:0141003 - p: 1.72e-05 *** histone kinase activity - GO:0035173 - p: 2.19e-05 *** histone H3 kinase activity - GO:0140996 - p: 1.47e-05 *** chromatin binding - GO:0003682 - p: 1.52e-05 *** 1.3 1.5 1.7 1.4 1.6 Absolute NES

GO Term