Enterocytes_TMIGD1_MEP1A - EPI - Colon - InactiveCD_vs_Control tube morphogenesis - GO:0035239 - p: 6.45e-10 *** regulation of anatomical structure morphogenesis - GO:0022603 - p: 4.63e-10 *** MAPK cascade - GO:0000165 - p: 1.05e-10 *** positive regulation of locomotion - GO:0040017 - p: 1.64e-10 *** positive regulation of cell motility - GO:2000147 - p: 3.26e-10 *** regulation of MAPK cascade - GO:0043408 - p: 2.78e-10 *** response to peptide hormone - GO:0043434 - p: 2.47e-10 *** vasculature development - GO:0001944 - p: 1.00e-10 *** blood vessel development - GO:0001568 - p: 1.00e-10 *** blood vessel morphogenesis - GO:0048514 - p: 1.00e-10 *** positive regulation of MAPK cascade - GO:0043410 - p: 7.16e-10 *** regulation of cytokine production - GO:0001817 - p: 3.25e-10 *** regulation of cell development - GO:0060284 - p: 1.62e-10 *** cytokine production - GO:0001816 - p: 2.03e-10 *** positive regulation of cell differentiation - GO:0045597 - p: 1.00e-10 *** positive regulation of immune response - GO:0050778 - p: 1.00e-10 *** 무 lymphocyte activation - GO:0046649 - p: 1.00e-10 *** Cluster immune effector process - GO:0002252 - p: 1.00e-10 *** T cell activation - GO:0042110 - p: 1.00e-10 *** cell-cell adhesion - GO:0098609 - p: 1.00e-10 *** regulation of lymphocyte activation - GO:0051249 - p: 1.00e-10 *** positive regulation of cell adhesion - GO:0045785 - p: 1.00e-10 *** GO Term regulation of cell-cell adhesion - GO:0022407 - p: 1.00e-10 *** regulation of leukocyte activation - GO:0002694 - p: 1.00e-10 *** Gene Count regulation of cell adhesion - GO:0030155 - p: 1.00e-10 *** positive regulation of cell activation - GO:0050867 - p: 9.25e-10 *** regulation of cell activation - GO:0050865 - p: 1.00e-10 *** positive regulation of cell-cell adhesion - GO:0022409 - p: 1.29e-10 *** regulation of T cell activation - GO:0050863 - p: 1.00e-10 *** regulation of leukocyte cell-cell adhesion - GO:1903037 - p: 1.00e-10 *** leukocyte cell-cell adhesion - GO:0007159 - p: 1.00e-10 *** adaptive immune response - GO:0002250 - p: 1.00e-10 *** positive regulation of leukocyte cell-cell adhesio... - GO:1903039 - p: 2.59e-10 *** transcription regulator complex - GO:0005667 - p: 6.97e-05 *** side of membrane - GO:0098552 - p: 4.22e-06 *** receptor complex - GO:0043235 - p: 3.17e-04 *** plasma membrane signaling receptor complex - GO:0098802 - p: 3.18e-04 *** kinase activity - GO:0016301 - p: 1.83e-05 *** phosphotransferase activity, alcohol group as acce... - GO:0016773 - p: 5.42e-07 *** histone modifying activity - GO:0140993 - p: 7.26e-10 *** protein kinase activity - GO:0004672 - p: 8.24e-09 *** histone H2AX kinase activity - GO:0141003 - p: 9.98e-07 *** histone kinase activity - GO:0035173 - p: 1.35e-07 *** histone H3 kinase activity - GO:0140996 - p: 1.86e-07 *** chromatin binding - GO:0003682 - p: 2.71e-05 *** 1.6 1.8 2.0 Absolute NES