Enterocytes_BEST4 - EPI - Ileum - InactiveCD_vs_Control cell-substrate adhesion - GO:0031589 - p: 1.56e-04 *** lymphocyte activation - GO:0046649 - p: 2.80e-04 *** cell activation - GO:0001775 - p: 1.55e-04 *** leukocyte activation - GO:0045321 - p: 2.40e-04 *** positive regulation of immune system process - GO:0002684 - p: 4.16e-05 *** T cell activation - GO:0042110 - p: 3.18e-04 *** cell adhesion - GO:0007155 - p: 1.25e-06 ** response to bacterium - GO:0009617 - p: 7.70e-05 *** cell-cell adhesion - GO:0098609 - p: 5.44e-06 *** inflammatory response - GO:0006954 - p: 1.03e-05 *** regulation of cell adhesion - GO:0030155 - p: 7.30e-06 *** regulation of leukocyte cell-cell adhesion - GO:1903037 - p: 1.10e-04 *** Gene Count positive regulation of T cell activation - GO:0050870 - p: 3.74e-04 *** leukocyte cell-cell adhesion - GO:0007159 - p: 8.17e-05 *** positive regulation of leukocyte cell-cell adhesio... - GO:1903039 - p: 1.41e-04 *** GO Term regulation of immune response - GO:0050776 - p: 1.31e-07 *** positive regulation of cell adhesion - GO:0045785 - p: 9.90e-06 ** Cluster positive regulation of cell-cell adhesion - GO:0022409 - p: 7.49e-05 *** innate immune response - GO:0045087 - p: 2.80e-08 *** adaptive immune response based on somatic recombin... - GO:0002460 - p: 4.66e-05 *** positive regulation of immune response - GO:0050778 - p: 1.04e-07 *** defense response to other organism - GO:0098542 - p: 2.73e-10 *** defense response to symbiont - GO:0140546 - p: 1.00e-10 *** adaptive immune response - GO:0002250 - p: 6.45e-07 *** membrane raft - GO:0045121 - p: 1.89e-03 * membrane microdomain - GO:0098857 - p: 2.18e-03 * endosome - GO:0005768 - p: 1.78e-04 *** membrane protein complex - GO:0098796 - p: 5.30e-05 *** side of membrane - GO:0098552 - p: 1.05e-05 ** plasma membrane signaling receptor complex - GO:0098802 - p: 1.27e-03 * plasma membrane protein complex - GO:0098797 - p: 1.73e-09 *** 1.6 1.8 2.0 Absolute NES