Enterocytes_TMIGD1_MEP1A - EPI - Ileum - InactiveCD_vs_Control response to type II interferon – GO:0034341 – p: 5.67e-07 *** positive regulation of cell motility - GO:2000147 - p: 2.36e-07 *** positive regulation of locomotion - GO:0040017 - p: 1.20e-07 *** regulation of hormone levels - GO:0010817 - p: 3.84e-09 *** positive regulation of immune response - GO:0050778 - p: 1.30e-07 *** regulation of leukocyte activation - GO:0002694 - p: 1.53e-07 *** positive regulation of cell adhesion - GO:0045785 - p: 6.02e-07 *** positive regulation of cytokine production - GO:0001819 - p: 9.10e-07 *** regulation of cytokine production - GO:0001817 - p: 3.49e-09 *** regulation of cell activation - GO:0050865 - p: 3.14e-08 *** Gene Count response to bacterium - GO:0009617 - p: 3.29e-08 *** cytokine production - GO:0001816 - p: 1.51e-09 *** positive regulation of leukocyte activation – GO:0002696 – p: 1.19e–06 ***

positive regulation of cell activation – GO:0050867 – p: 8.46e–07 *** Cluster regulation of leukocyte proliferation - GO:0070663 - p: 5.18e-07 *** regulation of inflammatory response - GO:0050727 - p: 5.88e-08 *** adaptive immune response - GO:0002250 - p: 3.05e-09 *** inflammatory response - GO:0006954 - p: 1.00e-10 *** plasma membrane protein complex - GO:0098797 - p: 3.73e-07 *** S side of membrane - GO:0098552 - p: 4.10e-09 *** cytokine receptor binding - GO:0005126 - p: 1.06e-03 ** signaling receptor activator activity - GO:0030546 - p: 7.03e-05 *** G protein-coupled receptor binding - GO:0001664 - p: 9.74e-05 *** signaling receptor regulator activity - GO:0030545 - p: 1.22e-05 *** chemokine receptor binding - GO:0042379 - p: 8.31e-05 *** 1.8 1.9 2.0 2.2 Absolute NES