Enterocytes_TMIGD1_MEP1A - EPI - Colon - ActiveCD_vs_Control lymphocyte activation - GO:0046649 - p: 1.00e-10 ** tube morphogenesis - GO:0035239 - p: 1.00e-10 ** positive regulation of immune response - GO:0050778 - p: 1.15e-10 *** regulation of immune response - GO:0050776 - p: 1.00e-10 ** positive regulation of cell population proliferati... - GO:0008284 - p: 1.00e-10 *** positive regulation of cell differentiation - GO:0045597 - p: 1.00e-10 ** regulation of cytokine production - GO:0001817 - p: 1.00e-10 *** regulation of cell activation - GO:0050865 - p: 4.00e-10 ** cytokine production - GO:0001816 - p: 1.00e-10 ** vasculature development - GO:0001944 - p: 1.00e-10 * blood vessel morphogenesis - GO:0048514 - p: 1.00e-10 ** regulation of cell-cell adhesion - GO:0022407 - p: 1.00e-10 ** blood vessel development - GO:0001568 - p: 1.00e-10 ** response to wounding – GO:0009611 – p: 1.00e–10 * inflammatory response - GO:0006954 - p: 1.00e-10 ** angiogenesis - GO:0001525 - p: 1.00e-10 ** Gene Count cell-cell adhesion - GO:0098609 - p: 1.00e-10 ** regulation of cell adhesion - GO:0030155 - p: 1.00e-10 ** wound healing - GO:0042060 - p: 1.00e-10 ** response to bacterium - GO:0009617 - p: 1.00e-10 ** positive regulation of cytokine production - GO:0001819 - p: 1.00e-10 ** positive regulation of cell migration - GO:0030335 - p: 1.00e-10 ** GO Term leukocyte cell-cell adhesion - GO:0007159 - p: 1.00e-10 ** Cluster positive regulation of cell adhesion - GO:0045785 - p: 1.00e-10 ** positive regulation of cell motility - GO:2000147 - p: 1.00e-10 ** positive regulation of locomotion - GO:0040017 - p: 1.00e-10 ** immune effector process - GO:0002252 - p: 1.00e-10 ** mononuclear cell proliferation - GO:0032943 - p: 3.57e-10 *** leukocyte proliferation - GO:0070661 - p: 3.66e-10 ** lymphocyte proliferation - GO:0046651 - p: 3.41e-10 *** adaptive immune response based on somatic recombin... - GO:0002460 - p: 2.42e-10 ** adaptive immune response - GO:0002250 - p: 1.00e-10 *** positive regulation of leukocyte proliferation - GO:0070665 - p: 5.39e-10 ** perinuclear region of cytoplasm - GO:0048471 - p: 2.98e-04 ** focal adhesion - GO:0005925 - p: 4.26e-04 * cell-substrate junction - GO:0030055 - p: 1.51e-04 ** plasma membrane protein complex - GO:0098797 - p: 6.70e-06 * side of membrane - GO:0098552 - p: 1.13e-08 * plasma membrane signaling receptor complex - GO:0098802 - p: 2.27e-05 * growth factor receptor binding - GO:0070851 - p: 5.02e-04 ** G protein-coupled receptor binding - GO:0001664 - p: 4.49e-04 ** ₹ chemokine receptor binding - GO:0042379 - p: 1.45e-03 cytokine receptor binding - GO:0005126 - p: 3.32e-06 ** cell adhesion molecule binding - GO:0050839 - p: 7.86e-06 * 1.7 1.9 1.3 Absolute NES