Fibroblasts_ADAMDEC1 - STR - Ileum - ActiveCD_vs_Control positive regulation of apoptotic process - GO:0043065 - p: 6.41e-07 *** positive regulation of programmed cell death - GO:0043068 - p: 9.45e-07 *** G protein-coupled receptor signaling pathway - GO:0007186 - p: 1.31e-07 *** inflammatory response - GO:0006954 - p: 1.05e-06 *** lymphocyte activation - GO:0046649 - p: 1.72e-07 *** regulation of cell activation - GO:0050865 - p: 1.54e-06 *** T cell activation - GO:0042110 - p: 3.56e-08 *** leukocyte cell-cell adhesion - GO:0007159 - p: 2.59e-07 *** regulation of cytokine production - GO:0001817 - p: 1.20e-10 *** cytokine production - GO:0001816 - p: 1.00e-10 *** positive regulation of cell-cell adhesion - GO:0022409 - p: 1.90e-06 *** regulation of cell-cell adhesion - GO:0022407 - p: 1.72e-08 *** regulation of cell adhesion - GO:0030155 - p: 1.00e-10 *** adaptive immune response - GO:0002250 - p: 1.55e-07 *** positive regulation of cytokine production - GO:0001819 - p: 1.59e-09 *** positive regulation of cell adhesion - GO:0045785 - p: 4.70e-10 *** MAPK cascade - GO:0000165 - p: 1.62e-06 *** ВР morphogenesis of an epithelium - GO:0002009 - p: 2.09e-06 *** regulation of MAPK cascade - GO:0043408 - p: 7.48e-07 *** tissue morphogenesis - GO:0048729 - p: 7.24e-08 *** mesenchyme development - GO:0060485 - p: 9.25e-07 *** Gene Count renal system development - GO:0072001 - p: 5.24e-07 *** angiogenesis - GO:0001525 - p: 2.70e-09 *** blood vessel morphogenesis - GO:0048514 - p: 1.00e-10 *** phosphatidylinositol 3-kinase/protein kinase B sig... - GO:0043491 - p: 2.88e-07 *** epithelial cell proliferation - GO:0050673 - p: 4.29e-09 *** regulation of epithelial cell proliferation - GO:0050678 - p: 7.03e-08 *** regulation of vasculature development - GO:1901342 - p: 2.04e-07 *** regulation of angiogenesis - GO:0045765 - p: 1.20e-07 *** regulation of phosphatidylinositol 3-kinase/protei... - GO:0051896 - p: 3.07e-07 *** Cluster blood vessel development - GO:0001568 - p: 1.00e-10 *** vasculature development - GO:0001944 - p: 1.00e-10 *** positive regulation of locomotion - GO:0040017 - p: 1.00e-10 *** positive regulation of cell motility - GO:2000147 - p: 1.00e-10 *** positive regulation of cell migration - GO:0030335 - p: 1.00e-10 *** cell-substrate junction - GO:0030055 - p: 9.86e-05 *** focal adhesion - GO:0005925 - p: 3.66e-05 *** CC plasma membrane protein complex - GO:0098797 - p: 8.22e-04 *** side of membrane - GO:0098552 - p: 9.24e-05 *** receptor complex - GO:0043235 - p: 1.19e-04 *** protein domain specific binding - GO:0019904 - p: 9.17e-04 *** histone modifying activity - GO:0140993 - p: 3.28e-04 *** kinase activity - GO:0016301 - p: 2.76e-05 *** phosphotransferase activity, alcohol group as acce... - GO:0016773 - p: 1.87e-05 *** protein kinase activity - GO:0004672 - p: 8.55e-06 *** histone H2AX kinase activity - GO:0141003 - p: 1.49e-04 *** histone H3 kinase activity - GO:0140996 - p: 3.42e-05 *** histone kinase activity - GO:0035173 - p: 8.00e-05 *** histone H2AXY142 kinase activity - GO:0140801 - p: 2.28e-03 ** histone H3Y41 kinase activity - GO:0035401 - p: 2.28e-03 ** protein tyrosine kinase activity - GO:0004713 - p: 1.49e-04 *** cell adhesion molecule binding - GO:0050839 - p: 1.05e-03 ** protein kinase binding - GO:0019901 - p: 1.12e-05 *** cytokine receptor binding - GO:0005126 - p: 9.22e-07 *** CCR chemokine receptor binding - GO:0048020 - p: 3.64e-05 *** signaling receptor activator activity - GO:0030546 - p: 2.22e-10 *** signaling receptor regulator activity - GO:0030545 - p: 1.00e-10 *** G protein-coupled receptor binding - GO:0001664 - p: 4.58e-10 *** 3 chemokine receptor binding - GO:0042379 - p: 7.21e-08 *** 1.6 1.8 2.0

Absolute NES

GO Term