Endothelial_cells_DARC - STR - Colon - ActiveCD_vs_Control vasculature development - GO:0001944 - p: 4.74e-06 *** blood vessel development - GO:0001568 - p: 2.63e-06 *** tube morphogenesis - GO:0035239 - p: 2.56e-06 *** epithelium development - GO:0060429 - p: 1.84e-07 *** animal organ morphogenesis - GO:0009887 - p: 1.72e-06 *** tube development - GO:0035295 - p: 3.78e-08 *** blood vessel morphogenesis - GO:0048514 - p: 2.80e-06 *** tissue morphogenesis - GO:0048729 - p: 9.93e-06 *** angiogenesis - GO:0001525 - p: 5.36e-06 *** ВР regulation of anatomical structure morphogenesis - GO:0022603 - p: 8.86e-08 *** regulation of cell adhesion - GO:0030155 - p: 6.34e-08 *** cell-cell adhesion - GO:0098609 - p: 6.55e-08 *** enzyme-linked receptor protein signaling pathway - GO:0007167 - p: 8.98e-07 *** cell surface receptor protein tyrosine kinase sign... - GO:0007169 - p: 1.29e-05 *** phosphorylation - GO:0016310 - p: 3.37e-08 *** response to nitrogen compound - GO:1901698 - p: 9.36e-09 *** Cluster protein phosphorylation - GO:0006468 - p: 2.17e-08 *** response to peptide hormone - GO:0043434 - p: 3.46e-06 *** cellular response to nitrogen compound - GO:1901699 - p: 3.71e-07 *** transcription regulator complex - GO:0005667 - p: 2.20e-03 ** GO Term perinuclear region of cytoplasm - GO:0048471 - p: 3.06e-05 *** somatodendritic compartment - GO:0036477 - p: 1.24e-04 *** Gene Count neuron projection - GO:0043005 - p: 2.18e-07 *** 100 dendritic tree - GO:0097447 - p: 1.13e-05 *** dendrite - GO:0030425 - p: 1.13e-05 *** axon - GO:0030424 - p: 1.38e-07 *** cell-substrate junction - GO:0030055 - p: 1.03e-04 *** focal adhesion - GO:0005925 - p: 1.03e-04 *** anchoring junction - GO:0070161 - p: 9.75e-09 *** protein phosphatase binding - GO:0019903 - p: 1.12e-05 *** transcription regulatory region nucleic acid bindi... - GO:0001067 - p: 2.22e-05 *** transcription cis-regulatory region binding - GO:0000976 - p: 2.22e-05 *** sequence-specific double-stranded DNA binding - GO:1990837 - p: 1.19e-06 *** chromatin binding - GO:0003682 - p: 8.47e-07 *** transferase activity, transferring phosphorus-cont... - GO:0016772 - p: 5.01e-06 *** kinase activity - GO:0016301 - p: 2.77e-07 *** protein kinase activity - GO:0004672 - p: 2.29e-06 *** phosphotransferase activity, alcohol group as acce... - GO:0016773 - p: 5.72e-08 *** histone modifying activity - GO:0140993 - p: 6.91e-09 *** histone kinase activity - GO:0035173 - p: 6.85e-10 *** histone H2AX kinase activity - GO:0141003 - p: 3.92e-10 *** histone H3 kinase activity - GO:0140996 - p: 4.47e-10 *** 1.3 1.5 1.7 1.4 1.6 Absolute NES