Endothelial_cells_CD36 - STR - Colon - ActiveCD_vs_Control regulation of cell adhesion - GO:0030155 - p: 1.12e-04 *** positive regulation of cell differentiation - GO:0045597 - p: 1.10e-04 *** regulation of anatomical structure morphogenesis – GO:0022603 – p: 7.05e–05 *** cell-cell adhesion - GO:0098609 - p: 1.08e-05 *** mesenchyme development - GO:0060485 - p: 2.33e-05 *** 0 perinuclear region of cytoplasm - GO:0048471 - p: 1.63e-03 ** neuron projection - GO:0043005 - p: 6.88e-06 *** somatodendritic compartment - GO:0036477 - p: 3.73e-05 *** 8 axon - GO:0030424 - p: 1.24e-04 *** Gene Count dendritic tree - GO:0097447 - p: 7.99e-06 *** dendrite - GO:0030425 - p: 7.99e-06 *** GO Term chromatin binding - GO:0003682 - p: 7.95e-05 *** Cluster histone binding - GO:0042393 - p: 2.67e-05 *** RNA polymerase II transcription regulatory region ... – GO:0000977 – p: 6.60e–05 *** DNA-binding transcription factor activity - GO:0003700 - p: 3.12e-05 *** DNA-binding transcription factor activity, RNA pol... - GO:0000981 - p: 7.05e-05 *** transcription regulatory region nucleic acid bindi... - GO:0001067 - p: 3.84e-06 *** transcription cis-regulatory region binding - GO:0000976 - p: 3.84e-06 *** kinase activity - GO:0016301 - p: 1.05e-04 *** phosphotransferase activity, alcohol group as acce... – GO:0016773 – p: 2.11e–05 *** protein kinase activity - GO:0004672 - p: 1.69e-05 *** histone modifying activity - GO:0140993 - p: 1.72e-07 *** histone kinase activity - GO:0035173 - p: 1.29e-05 *** histone H2AX kinase activity - GO:0141003 - p: 1.09e-05 *** histone H3 kinase activity - GO:0140996 - p: 1.04e-05 *** 1.3 1.5 Absolute NES