Endothelial_cells_CD36 - STR - Colon - InactiveCD_vs_Control cellular response to hormone stimulus – GO:0032870 – p: 3.62e–05 *** cell surface receptor protein tyrosine kinase sign... - GO:0007169 - p: 2.44e-05 *** response to peptide hormone - GO:0043434 - p: 5.81e-05 *** cellular response to peptide hormone stimulus – GO:0071375 – p: 5.30e–05 *** peptidyl-amino acid modification - GO:0018193 - p: 5.98e-06 *** positive regulation of cell differentiation – GO:0045597 – p: 8.82e–06 *** regulation of anatomical structure morphogenesis – GO:0022603 – p: 2.22e–07 *** axon development - GO:0061564 - p: 5.02e-05 *** transcription regulator complex - GO:0005667 - p: 5.36e-04 *** Gene Count receptor complex - GO:0043235 - p: 1.30e-04 *** somatodendritic compartment - GO:0036477 - p: 3.58e-04 *** GO Term dendritic tree - GO:0097447 - p: 6.18e-04 *** Cluster dendrite - GO:0030425 - p: 6.18e-04 *** postsynapse - GO:0098794 - p: 9.08e-05 *** axon - GO:0030424 - p: 1.72e-05 *** chromatin binding - GO:0003682 - p: 1.96e-08 *** histone binding - GO:0042393 - p: 2.69e-07 *** kinase activity - GO:0016301 - p: 1.99e-08 *** phosphotransferase activity, alcohol group as acce... - GO:0016773 - p: 2.57e-09 *** protein kinase activity - GO:0004672 - p: 2.11e-10 *** histone modifying activity - GO:0140993 - p: 1.00e-10 *** histone kinase activity - GO:0035173 - p: 2.61e-08 *** histone H3 kinase activity - GO:0140996 - p: 4.83e-08 *** histone H2AX kinase activity - GO:0141003 - p: 1.93e-08 *** 1.3 1.5 1.6 Absolute NES