Myofibroblasts_HHIP_NPNT - STR - Colon - InactiveCD_vs_Control tube morphogenesis - GO:0035239 - p: 1.57e-06 *** animal organ morphogenesis - GO:0009887 - p: 1.43e-06 *** regulation of cell adhesion - GO:0030155 - p: 1.69e-06 *** negative regulation of transcription by RNA polyme... - GO:0000122 - p: 1.09e-08 *** response to hormone - GO:0009725 - p: 8.05e-07 *** enzyme-linked receptor protein signaling pathway - GO:0007167 - p: 6.20e-09 *** cell surface receptor protein tyrosine kinase sign... - GO:0007169 - p: 5.32e-07 *** cellular response to hormone stimulus - GO:0032870 - p: 4.13e-07 *** response to peptide hormone - GO:0043434 - p: 1.67e-06 *** cellular response to nitrogen compound - GO:1901699 - p: 6.03e-08 *** phosphorylation - GO:0016310 - p: 1.00e-10 *** protein phosphorylation - GO:0006468 - p: 1.00e-10 *** neuron projection - GO:0043005 - p: 5.07e-07 *** Cluster somatodendritic compartment - GO:0036477 - p: 8.13e-06 *** dendritic tree - GO:0097447 - p: 1.40e-05 *** dendrite - GO:0030425 - p: 1.40e-05 *** GO Term axon - GO:0030424 - p: 4.94e-06 ** anchoring junction - GO:0070161 - p: 4.81e-07 *** Gene Count transcription regulator complex - GO:0005667 - p: 1.97e-05 *** chromatin binding - GO:0003682 - p: 2.45e-09 *** histone binding - GO:0042393 - p: 5.75e-07 *** cis-regulatory region sequence-specific DNA bindin... - GO:0000987 - p: 5.58e-10 *** RNA polymerase II transcription regulatory region ... - GO:0000977 - p: 1.38e-10 *** RNA polymerase II cis-regulatory region sequence-s... - GO:0000978 - p: 7.32e-10 *** DNA-binding transcription factor activity - GO:0003700 - p: 1.00e-10 *** DNA-binding transcription factor activity, RNA pol... - GO:0000981 - p: 1.00e-10 *** transferase activity, transferring phosphorus-cont... - GO:0016772 - p: 8.27e-06 *** kinase activity - GO:0016301 - p: 1.49e-07 *** phosphotransferase activity, alcohol group as acce... - GO:0016773 - p: 2.30e-08 *** protein kinase activity - GO:0004672 - p: 1.11e-10 *** histone modifying activity - GO:0140993 - p: 1.00e-10 *** histone H2AX kinase activity - GO:0141003 - p: 1.00e-10 ** histone H3 kinase activity - GO:0140996 - p: 1.00e-10 *** histone kinase activity - GO:0035173 - p: 1.00e-10 *** 1.4 1.5 1.6 1.7 Absolute NES