Fibroblasts_ADAMDEC1 - STR - Colon - InactiveCD_vs_Control actin cytoskeleton organization - GO:0030036 - p: 1.16e-06 *** anterograde trans-synaptic signaling - GO:0098916 - p: 1.78e-07 *** chemical synaptic transmission - GO:0007268 - p: 1.78e-07 *** synaptic signaling - GO:0099536 - p: 1.99e-07 *** trans-synaptic signaling - GO:0099537 - p: 9.28e-08 *** axon development - GO:0061564 - p: 3.94e-08 *** MAPK cascade - GO:0000165 - p: 2.03e-07 *** cellular response to nitrogen compound – GO:1901699 – p: 1.19e-07 *** regulation of protein phosphorylation - GO:0001932 - p: 1.97e-07 *** regulation of phosphorylation - GO:0042325 - p: 5.14e-08 *** cell surface receptor protein tyrosine kinase sign... - GO:0007169 - p: 6.52e-09 *** regulation of MAPK cascade - GO:0043408 - p: 1.31e-08 *** positive regulation of MAPK cascade - GO:0043410 - p: 7.45e-07 *** cellular response to peptide hormone stimulus - GO:0071375 - p: 3.20e-07 *** Cluster protein autophosphorylation - GO:0046777 - p: 1.04e-07 *** vasculature development - GO:0001944 - p: 2.99e-09 *** blood vessel development - GO:0001568 - p: 5.69e-09 *** blood vessel morphogenesis - GO:0048514 - p: 5.69e-09 *** tissue morphogenesis - GO:0048729 - p: 1.39e-08 *** GO Term receptor complex - GO:0043235 - p: 3.30e-04 *** transcription regulator complex - GO:0005667 - p: 1.71e-04 *** glutamatergic synapse - GO:0098978 - p: 1.05e-05 *** Gene Count axon - GO:0030424 - p: 1.53e-06 *** somatodendritic compartment - GO:0036477 - p: 2.57e-09 *** postsynapse - GO:0098794 - p: 1.00e-10 *** dendritic tree - GO:0097447 - p: 1.60e-09 *** dendrite - GO:0030425 - p: 1.60e-09 *** protein domain specific binding - GO:0019904 - p: 4.67e-06 *** protein kinase binding - GO:0019901 - p: 3.46e-06 *** chromatin binding - GO:0003682 - p: 4.77e-08 *** histone binding - GO:0042393 - p: 4.97e-05 *** transcription factor binding - GO:0008134 - p: 1.77e-05 *** kinase activity - GO:0016301 - p: 1.00e-10 *** phosphotransferase activity, alcohol group as acce... - GO:0016773 - p: 1.00e-10 *** histone modifying activity - GO:0140993 - p: 1.00e-10 *** protein kinase activity - GO:0004672 - p: 1.00e-10 *** histone kinase activity - GO:0035173 - p: 1.00e-10 *** histone H3 kinase activity - GO:0140996 - p: 1.00e-10 *** histone H2AX kinase activity - GO:0141003 - p: 1.00e-10 *** 1.3 1.4 1.5 Absolute NES