IncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns (23/945) = (func/total IncRNA) ::: biotype = lincRNA PageRank, range = (-0.1, 1.1)2.0 1.5 factor(label) density 0.1 0.5 -0.0 -0.4 8.0 0.0 rank