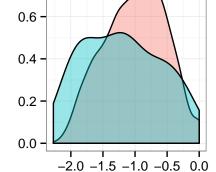lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns
(48 / 2413) = (func/total lncRNA) ::: biotype = lincProcTrans
measures across all samples(cell−type/pulldown)