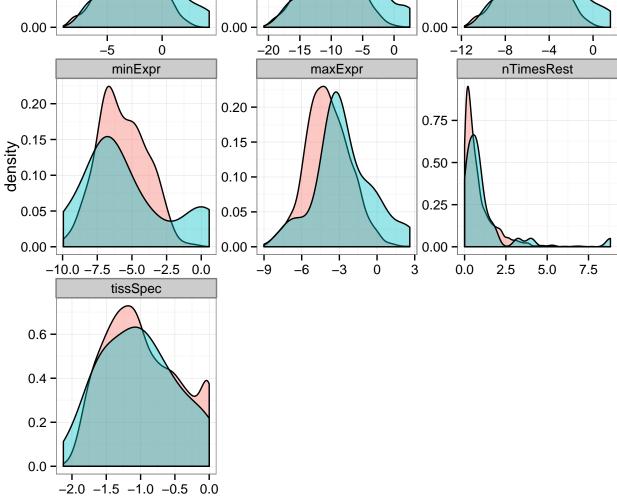lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns
(31 / 1889) = (func/total lncRNA) ::: biotype = antisense
measures across all samples(cell–type/pulldown)