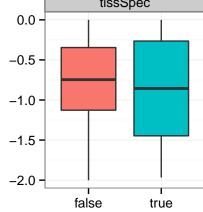lncRNA group = IDRlessthan0_2 ::: data cols = lpa
(31 / 2051) = (func/total lncRNA) ::: biotype = lincRNA
measures across all samples(cell−type/pulldown)