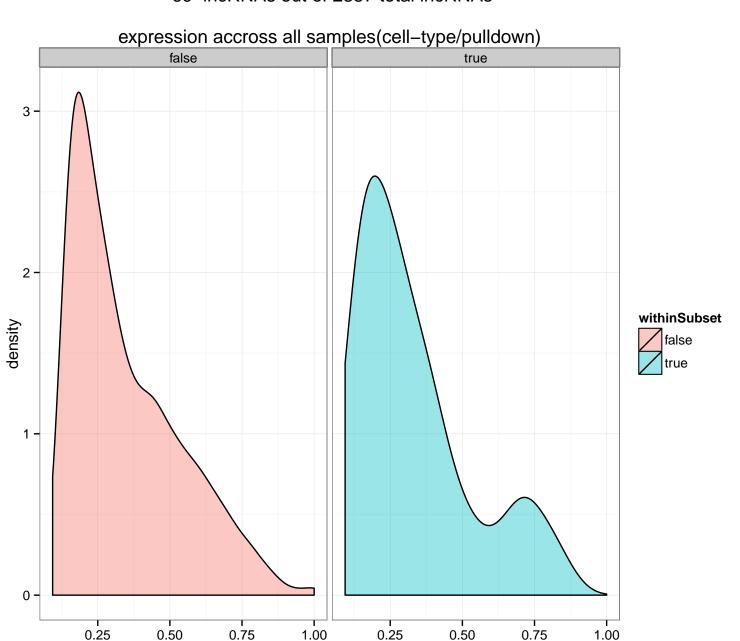
IncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (69 / 2857) = (func/total IncRNA) ::: biotype = all_biotypes subset = 69 IncRNAs out of 2857 total IncRNAs



JSD for each gene