IncRNA group = IDRlessthan0\_01 ::: data cols = lpa (4 / 72) = (func/total IncRNA) ::: biotype = antisense expression accross all samples(cell-type/pulldown) false 2.5 2.0 1.5 withinSubset density false true 1.0 -0.5 -0.0 -0.25 0.50 0.75 1.00 0.25 0.50 0.75 1.00

JSD for each gene