IncRNA group = IDRIessthan0_1 ::: data cols = bothPullDowns (46 / 1955) = (func/total IncRNA) ::: biotype = remove_antisense subset = 46 IncRNAs out of 1955 total IncRNAs average expression vs. tissue specificity (JSD) 1.00 -0.75 factor(withinSubset) false • true label • 0.00 • 0.25 • 0.50 0.75 0.50 -1.00 0.25 -

log(averageExpr)

-6