IncRNA group = IDRlessthan0_1 ::: data cols = Ipa (45 / 1554) = (func/total IncRNA) ::: biotype = remove_antisense average expression vs. tissue specificity (JSD) 1.00 -0.75 tissSpec factor(withinSubset) • false • true 0.50 -0.25 -15 10 averageExpr