IncRNA group = IDRlessthan0_01 ::: data cols = Inpa (21 / 146) = (func/total IncRNA) ::: biotype = lincProcTrans average expression vs. tissue specificity (JSD) 0.75 factor(withinSubset) • false • true 0.25 -2.5 7.5 0.0 5.0 averageExpr