IncRNA group = IDRIessthan0_1 ::: data cols = Inpa (42 / 1350) = (func/total IncRNA) ::: biotype = lincProcTrans average expression vs. tissue specificity (JSD) 0.75 tissSpec factor(withinSubset) • false • true 0.50 -0.25 -0.0 2.5 7.5 5.0 averageExpr