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 tissSpec 0.50 label • 0.00 • 0.25 • 0.50 0.75 1.00 0.25 --2.5 0.0 2.5 log(averageExpr)