IncRNA group = IDRlessthan0\_01 ::: data cols = Inpa ( 25 / 242 ) = (func/total IncRNA) ::: biotype = all\_biotypes 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)