IncRNA group = IDRlessthan0\_01 ::: data cols = bothPullDowns (25 / 243) = (func/total IncRNA) ::: biotype = all\_biotypes average expression vs. tissue specificity (JSD) 1.00 -0.75 tissSpec factor(withinSubset) • false • true 0.50 -0.25 -2.5 7.5 0.0 5.0 averageExpr