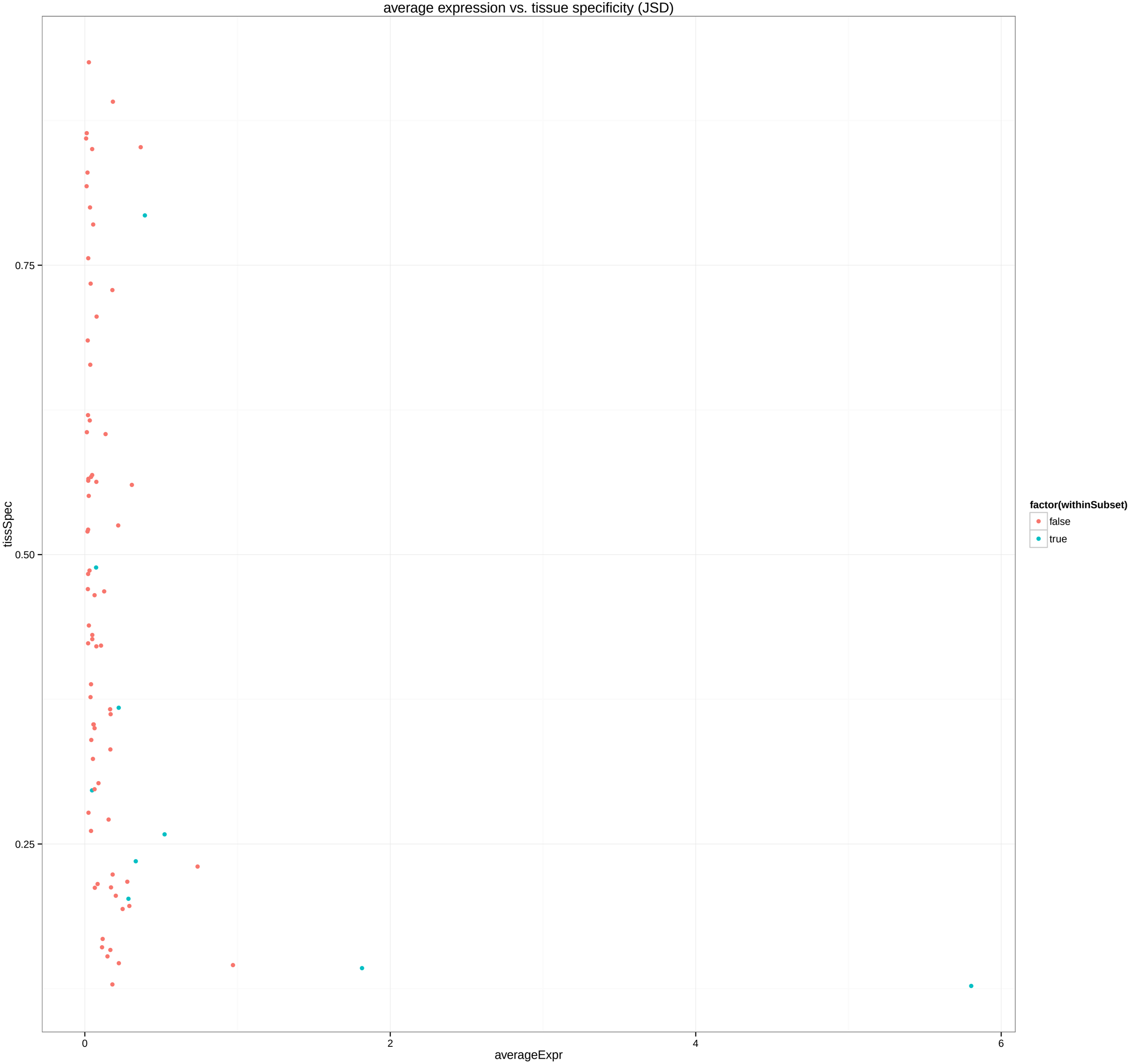
lncRNA group = IDRlessthan0_01 ::: data cols = lnpa
(9 / 81) = (func/total lncRNA) ::: biotype = lincRNA
subset =
9 lncRNAs out of 81 total lncRNAs

average expression vs. tissue specificity (JSD)