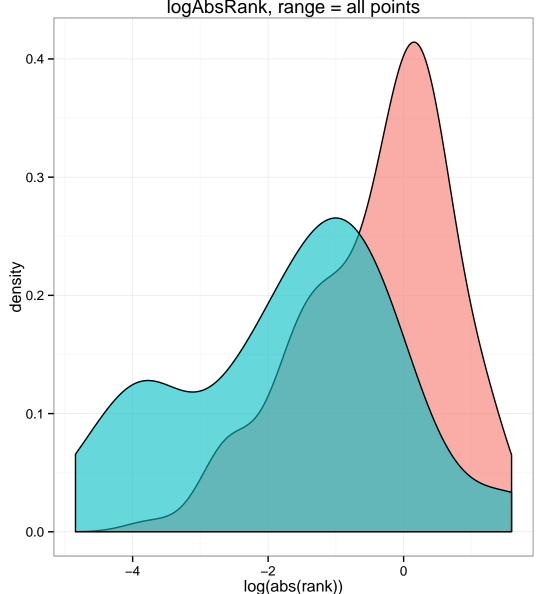
lncRNA group = IDRlessthan0_01 ::: data cols = Inpa (21 / 146) = (func/total IncRNA) ::: biotype = lincProcTrans logAbsRank, range = all points



factor(label)