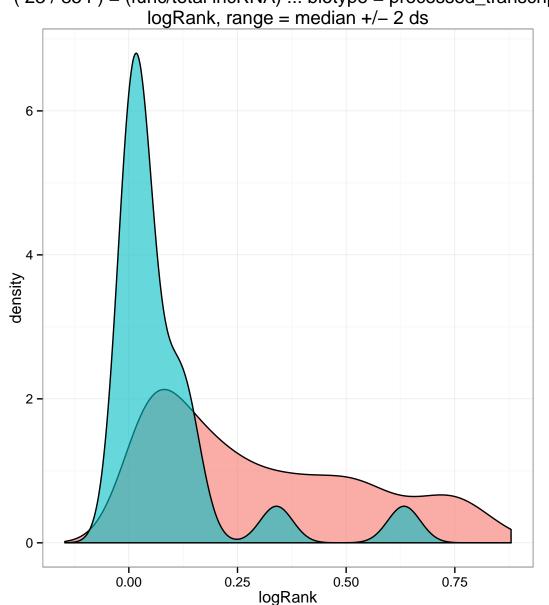
IncRNA group = IDRlessthan0\_2 ::: data cols = Inpa ( 23 / 854 ) = (func/total IncRNA) ::: biotype = processed\_transcript



factor(label)