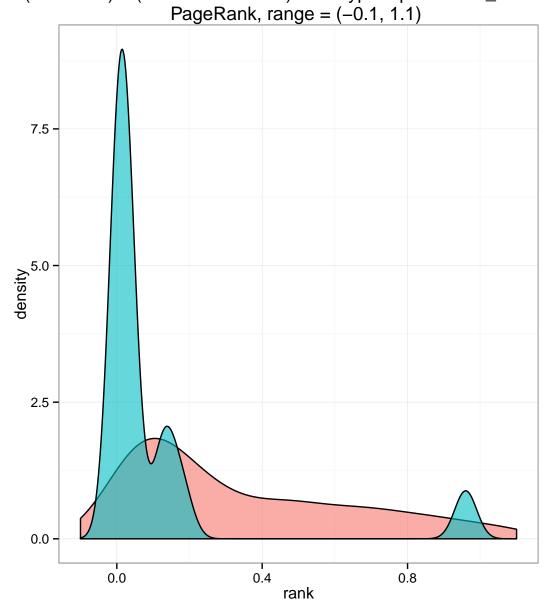
IncRNA group = IDRlessthan0\_2 ::: data cols = Inpa ( 17 / 328 ) = (func/total IncRNA) ::: biotype = processed\_transcript PageRank, range = (-0.1, 1.1)



factor(label)