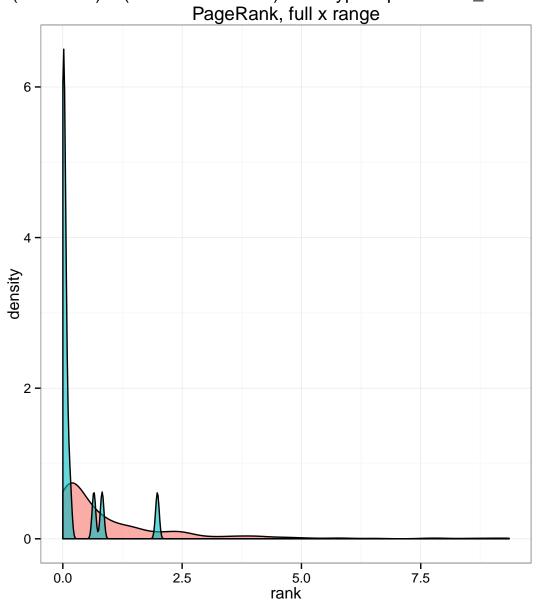
IncRNA group = IDRIessthan0_2 ::: data cols = bothPullDowns (17 / 329) = (func/total IncRNA) ::: biotype = processed_transcript PageRank, full x range



factor(label)