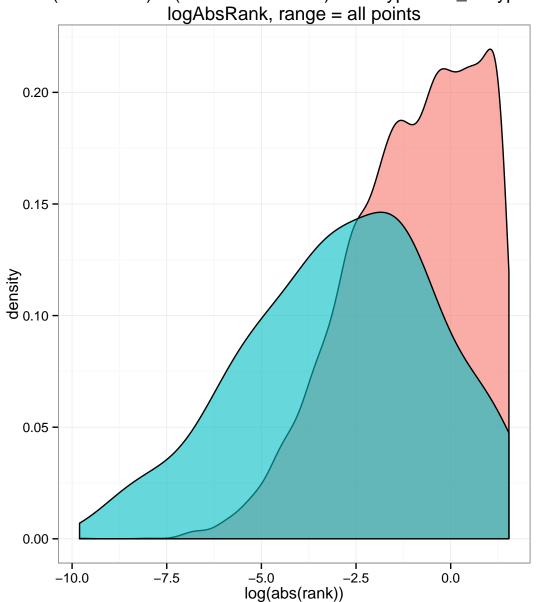
IncRNA group = allLncRNA ::: data cols = Inpa (87 / 6320) = (func/total IncRNA) ::: biotype = all_biotypes logAbsRank, range = all points



factor(label)