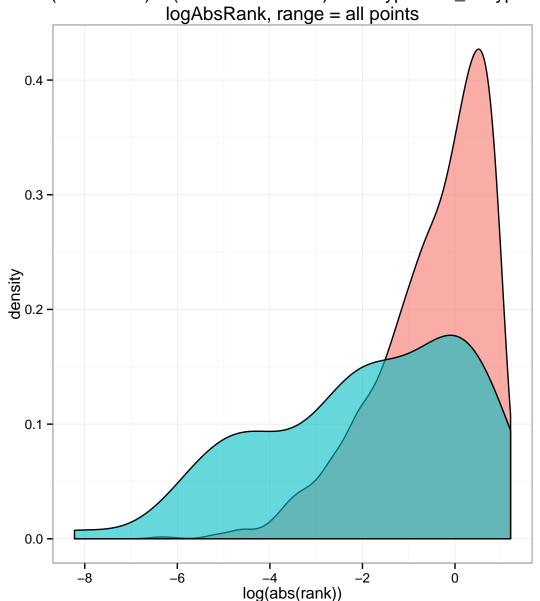
lncRNA group = IDRlessthan0_1 ::: data cols = lpa (69 / 2848) = (func/total lncRNA) ::: biotype = all_biotypes logAbsRank, range = all points



factor(label)