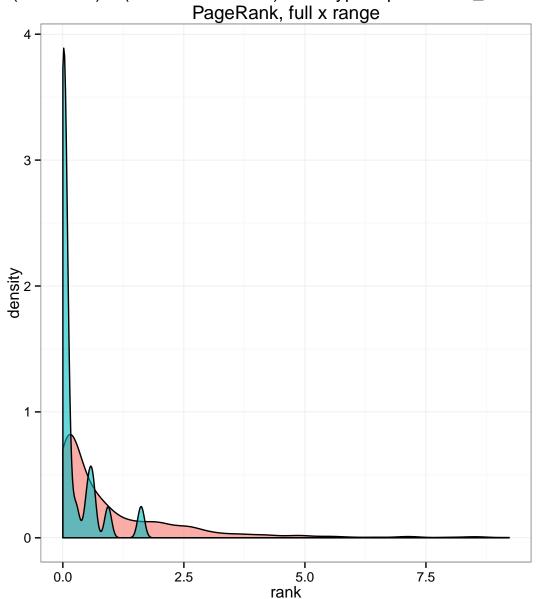
IncRNA group = IDRlessthan0_2 ::: data cols = lpa
(23 / 858) = (func/total IncRNA) ::: biotype = processed_transcript



factor(label)