IncRNA group = IDRlessthan0\_2 ::: data cols = lpa (82 / 5045) = (func/total IncRNA) ::: biotype = all\_biotypes PageRank, range = median +/- 2 ds 1000 -750 -500 -250 factor(label) 0 0 40 30 20 10 0 -0

rank

count